

**The action of xanomeline on human muscarinic receptors  
expressed in Chinese hamster ovary cells**

A DISSERTATION  
SUBMITTED TO THE FACULTY OF THE GRADUATE SCHOOL  
OF THE UNIVERSITY OF MINNESOTA  
BY

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IN PARTIAL FULFILLMENT OF THE REQUIREMENTS  
FOR THE DEGREE OF  
DOCTOR OF PHILOSOPHY

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**JANUARY, 2009**

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## **Acknowledgements**

I would like to thank my advisor Dr. Esam El-Fakahany for his support and guidance. He kindly let me join his lab and be his last graduate student. He has also helped me to grow as a scientist.

I would like to thank my committee members Dr. Virginia Seybold, Dr. William Engeland and Dr. Ping Law for their time and advice. Without their thoughtful input I would not have been able to complete my thesis as quickly. They have also been instrumental in helping me learn to present my work in a clear and concise manner.

The El-Fakahany lab members that I have worked with have been great. I would especially like to thank Marianne Grant for all of her help and support. She was always there to teach me new techniques, answer questions and help with anything. Without her I would not have been able to get everything done. I would also like to thank Kayla De Lorme and Allison Cherry for the great conversions and making the lab a fun place to be. I would also like to acknowledge both Marianne Grant and Kayla De Lorme for providing some of the data presented in this thesis.

I would also like to thank my former lab. Dr. Walter Low has been a great colleague over the years. I would specifically like to thank Terry Burns for his help and thoughtful discussions. I would also like to thank Aleta Rees, David Horn and Yuliya Perepelitsa for their hard work and helping me learn to be a better mentor.

Lastly, I would like to thank my friends and family for supporting me through all of my schooling. They have always been there when I needed them, in both the good times and the bad. I would especially like to thank my parents for their support and encouragement over the years. I would like to thank my boyfriend, Mark Gramith, who has helped me through each stage and has had an incredible amount of patience. I

would also like to thank Thaddeus Brink for taking the time to read through my thesis and giving my feedback.

*This dissertation is dedicated to my grandfather David Noetzel,  
who got me interested in science at a very young age.*

## **Thesis abstract**

Muscarinic acetylcholine receptors are part of the G protein-coupled superfamily of receptors. There are five subtypes of muscarinic receptors (M1-M5). These receptors are therapeutic targets for in a number of diseases, including Alzheimer's disease and schizophrenia. The orthosteric (primary) binding domain of muscarinic receptors is highly conserved across subtypes making it difficult to develop agonists or antagonists that bind with selectivity at a particular subtype. However, there are allosteric (secondary) binding sites on muscarinic receptors that are thought to be less well conserved between subtypes, which may lead to the development of drugs that bind in a subtype-selective manner. One potential drug of interest is xanomeline. Previous research has found that it binds in a unique manner to muscarinic receptors; it binds in a reversible manner at the orthosteric site and in a wash-resistant manner at an allosteric site. Furthermore, xanomeline is thought to be an M1/M4 selective agonist. The goal of the current research was to characterize the effects of prolonged exposure of the various receptor subtypes to xanomeline and determine the mechanisms through which these effects occur. To do this Chinese hamster ovary cells expressing the various individual muscarinic receptors were exposed to xanomeline for brief and long-term time periods and radioligand binding and functional assays were performed. Xanomeline was shown to bind in a reversible and wash-resistant manner at the M1-M4 receptor subtypes. Although the effects of acute xanomeline exposure were similar across subtypes, long-term treatment with xanomeline resulted in differential effects among subtypes. More detailed experiments were conducted at the M1 and M3 receptor subtypes to elucidate the mechanisms of the long-term effects of xanomeline. Wash-resistant xanomeline binding was able to modulate the functional properties of the receptor. The consequences of the long-term effects of wash-resistant xanomeline binding are

dependent on activation of the receptor through the orthosteric binding site. It appears that these effects are a result of receptor internalization and allosteric modulation of the receptor. Thus, my research demonstrates that it is important to determine both the acute and long-term drug interactions with the receptor as part of the drug development process.

## **TABLE OF CONTENTS**

<b>Chapter 1: Introduction – muscarinic receptors and xanomeline</b>	
1. Overview	2
2. G protein-coupled receptors	4
3. Subtypes, localization and function of muscarinic receptors	5
4. Structure and binding of muscarinic receptors	7
5. A model of receptor activation	9
6. Receptor trafficking	10
7. Drug-receptor interactions	10
8. Signaling pathways for muscarinic receptors	11
9. Consequences of prolonged exposure of muscarinic receptors to agonist and antagonists	15
10. Influence of receptor mutations and expression levels on receptor activity	20
11. In vitro receptor expression systems	21
12. Allosteric interactions with the receptor	23
13. Pharmacological profile of allosteric binding	24
14. Allosteric sites on muscarinic receptors	25
15. Muscarinic receptors, allosteric sites and diseases	28
16. Alzheimer’s disease	28
17. Cholinergic hypothesis of Alzheimer’s disease	29
18. Cholinergic treatments for Alzheimer’s disease	31
19. Non-cholinergic treatments for Alzheimer’s disease	32
20. Schizophrenia	33
21. Treatment for schizophrenia	35
22. Binding and functional properties of xanomeline	35
23. Proposed mechanism of xanomeline binding	37
24. Wash-resistant xanomeline binding	38
25. Xanomeline as a treatment for Alzheimer’s disease and schizophrenia	39
26. Xanomeline treatment in Alzheimer’s disease models	40
27. Clinical trials with xanomeline for the treatment of Alzheimer’s disease	41
28. Xanomeline treatments in animal models of schizophrenia	42
29. Clinical trials with xanomeline for the treatment of Alzheimer’s disease	43
30. Summary	43
<b>Chapter 2: Materials and methods</b>	
1. Materials and general treatment protocols	46
2. Radioligand binding assays	48
3. Functional assays	50
4. Variations on radioligand binding and functional assay paradigms	54
5. Data analysis	59
<b>Chapter 3: Determination of the acute and long-term effects of xanomeline binding at M1-M4 muscarinic receptors</b>	
1. Introduction	61
2. Results	62
3. Discussion	72
4. Figures	80
5. Tables	115



<b>Chapter 4: Determination of the acute and long-term functional effects of xanomeline at the M1 and M3 muscarinic receptors</b>	
1. Introduction	121
2. Results	122
3. Discussion	131
4. Figures	136
5. Tables	152
<b>Chapter 5: Receptor activation is required for the development of the long-term effects of xanomeline at the M1 and M3 muscarinic receptors</b>	
1. Introduction	158
2. Results	159
3. Discussion	167
4. Figures	172
5. Tables	188
<b>Chapter 6: Mechanisms of the long-term effects of xanomeline wash-resistant binding</b>	
1. Introduction	196
2. Results	197
3. Discussion	204
4. Figures	210
5. Tables	219
<b>Chapter 7: Conclusions</b>	224
<b>References</b>	230
<b>Appendix A: Differences observed between suspension and monolayer protocols</b>	
1. Introduction	242
2. Results	242
3. Discussion	244
4. Figures	246
5. Table	248

## **LIST OF TABLES**

### **Chapter 3: Determination of the acute and long-term effects of xanomeline binding at M1-M4 muscarinic receptors**

Table 1. Short- and long-term effects of xanomeline, carbachol or pilocarpine binding on [ <sup>3</sup> H]NMS binding in CHO cells expressing hM1-hM4 receptor subtypes.	115
Table 2. [ <sup>3</sup> H]NMS saturation binding parameters for xanomeline, carbachol or pilocarpine pretreatment and waiting periods in CHO cells expressing hM1-hM4 receptor subtypes.	118

### **Chapter 4: Determination of the acute and long-term functional effects of xanomeline at the M1 and M3 muscarinic receptors**

Table 1. Effects of agonist treatment on PI hydrolysis in CHO cells expressing hM1 and hM3 receptor subtypes.	152
Table 2. Effects of carbachol, pilocarpine or xanomeline stimulation following short- and long-term agonist pretreatment and washout on inositol phosphates production in CHO cells expressing hM3 receptors.	153
Table 3. Effects of carbachol, pilocarpine or xanomeline stimulation following short- and long-term agonist pretreatment and washout on inositol phosphates production in CHO cells expressing hM1 receptors.	155

### **Chapter 5: Receptor activation is required for the development of the long-term effects of xanomeline at the M1 and M3 muscarinic receptors**

Table 1. Effects of xanomeline pretreatment and washout in the absence or presence of atropine on [ <sup>3</sup> H]NMS saturation binding and functional parameters of carbachol stimulation in CHO cells expressing hM1 or hM3 receptors.	188
Table 2. Effects of xanomeline pretreatment on inhibition of NMS binding, saturation binding and functional parameters in wild-type and mutant rM1 receptors expressed in CHO cells.	190
Table 3. Effects of blocking receptor function using siRNA against G <sub>q</sub> and G <sub>11</sub> G 192 proteins on [ <sup>3</sup> H]NMS saturation binding parameters in CHO cells expressing hM3 receptors assayed in suspension.	
Table 4. Effects of xanomeline analogs on inhibition of [ <sup>3</sup> H]NMS binding and functional parameters in CHO cells expressing hM3 receptors.	193
Table 5. Effects of treatment with the propyl xanomeline analog on [ <sup>3</sup> H]NMS saturation binding parameters in CHO cells expressing hM3 receptors.	194

## **Chapter 6: Mechanisms of the long-term effects of xanomeline wash-resistant binding**

- Table 1. Inhibition of radioligand binding using a receptor-saturating concentration of [<sup>3</sup>H]QNB or [<sup>3</sup>H]NMS following xanomeline pretreatment and various waiting periods in CHO cells expressing hM1 or hM3 receptor subtypes. 219
- Table 2. Effects of blocking receptor activation on xanomeline-induced changes in [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing hM3 receptors. 221
- Table 3. Effects of changing the level of receptor expression or inactivating the G proteins on xanomeline-induced changes in [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing the hM4 receptor subtype. 222

## **Appendix A: Differences observed between suspension and monolayer protocols**

- Table 1. Effects on [<sup>3</sup>H]NMS saturation binding parameters when CHO hM3 cells were assayed in monolayer versus suspension. 248

## **LIST OF FIGURES**

### **Chapter 3: Determination of the acute and long-term effects of xanomeline binding at M1-M4 muscarinic receptors**

Figure 1.	Time course of the development of xanomeline wash-resistant binding at hM1-hM4 muscarinic receptor subtypes expressed in CHO cells.	80
Figure 2.	Inhibition of [ <sup>3</sup> H]NMS binding for acute xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	82
Figure 3.	Effects of acute DMSO pretreatments on the inhibition of [ <sup>3</sup> H]NMS binding in CHO cells stably expressing hM1-hM4 receptor subtypes.	84
Figure 4.	Effects of acute xanomeline pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	86
Figure 5.	Effects of acute DMSO pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	88
Figure 6.	Inhibition of [ <sup>3</sup> H]NMS binding for acute carbachol pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	90
Figure 7.	Effects of acute carbachol pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	92
Figure 8.	Inhibition of [ <sup>3</sup> H]NMS binding for acute pilocarpine pretreatments in CHO cells stably expressing the hM3 receptor subtype.	94
Figure 9.	Effects of acute pilocarpine pretreatments on protein content in CHO cells stably expressing the hM3 receptor subtype.	95
Figure 10.	[ <sup>3</sup> H]NMS saturation binding subsequent to acute xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	96
Figure 11.	Inhibition of [ <sup>3</sup> H]NMS binding for long-term xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	98
Figure 12.	Effects of long-term DMSO pretreatments on the inhibition of [ <sup>3</sup> H]NMS binding in CHO cells stably expressing hM1-hM4 receptor subtypes.	100
Figure 13.	Effects of long-term xanomeline pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	102
Figure 14.	Effects of long-term DMSO pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	104
Figure 15.	Inhibition of [ <sup>3</sup> H]NMS binding for long-term carbachol pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	106
Figure 16.	Effects of long-term carbachol pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.	108
Figure 17.	Inhibition of [ <sup>3</sup> H]NMS binding for long-term pilocarpine pretreatments in CHO cells stably expressing the hM3 receptor subtype.	110
Figure 18.	Effects of long-term pilocarpine pretreatments on protein content in CHO cells stably expressing hM3 receptor subtype.	111

Figure 19.	[ <sup>3</sup> H]NMS saturation binding subsequent to long-term xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.	112
Figure 20.	[ <sup>3</sup> H]NMS saturation binding subsequent to long-term agonist pretreatments in CHO cells stably expressing the hM3 receptor subtype.	114

#### **Chapter 4: Determination of the acute and long-term functional effects of xanomeline at the M1 and M3 muscarinic receptors**

Figure 1.	hM3 receptor activation by xanomeline reversible and wash-resistant binding.	136
Figure 2.	hM1 receptor activation by xanomeline reversible and wash-resistant binding.	137
Figure 3.	Effects of acute xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM3 receptor.	138
Figure 4.	Effects of acute xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM1 receptor.	140
Figure 5.	Effects of acute pilocarpine on the ability of pilocarpine to subsequently stimulate the production of inositol phosphates at the hM3 receptor.	142
Figure 6.	Stimulation of inositol phosphate production by exogenous phospholipase C following brief xanomeline pretreatments in cells expressing hM1 and hM3 receptors.	143
Figure 7.	Stimulation of inositol phosphate production by exogenous phospholipase C following long-term xanomeline pretreatments in cells expressing hM1 and hM3 receptors.	144
Figure 8.	Effects of prolonged xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM3 receptor.	145
Figure 9.	Effects of prolonged xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM1 receptor.	147
Figure 10.	Effects of long-term agonist exposure on the ability of various muscarinic agonists to stimulate inositol phosphates production at the hM1 and hM3 receptors.	149
Figure 11.	Effects of long-term exposure to carbachol on the ability of xanomeline to stimulate inositol phosphate production at hM1 and hM3 receptors.	151

#### **Chapter 5: Receptor activation is required for the development of the long-term effects of xanomeline at the M1 and M3 muscarinic receptors**

Figure 1.	Effects of atropine on long-term xanomeline-induced inhibition of [ <sup>3</sup> H]NMS binding in CHO cells expressing hM3 receptors.	172
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Figure 2.	Effects of atropine on long-term xanomeline-induced inhibition of [ <sup>3</sup> H]NMS binding in CHO cells expressing hM1 receptors.	174
Figure 3.	Effects of atropine on long-term modulation of inositol phosphates production by xanomeline in CHO cells expressing hM3 receptors.	176
Figure 4.	Effects of atropine on long-term modulation of inositol phosphates production by xanomeline in CHO cells expressing hM1 receptors.	178
Figure 5.	Effects of using a non-functional receptor mutant on function, inhibition of [ <sup>3</sup> H]NMS and saturation binding in CHO cells expressing rM1 receptors.	180
Figure 6.	Control experiments to determine [ <sup>3</sup> H]NMS binding and protein content following xanomeline or DMSO pretreatments in CHO cells expressing rM1 receptors.	182
Figure 7.	Effects of blocking receptor function using siRNA against G <sub>q</sub> and G <sub>11</sub> G proteins on long-term xanomeline-induced inhibition of [ <sup>3</sup> H]NMS binding in CHO cells expressing the hM3 receptor.	184
Figure 8.	Effects of wash-resistant binding in the absence of receptor activation on the ability of the propyl analog of xanomeline to initiate long-term effects on saturation binding parameters in CHO cells expressing hM3 receptors.	186

## **Chapter 6: Mechanisms of the long-term effects of xanomeline wash-resistant binding**

Figure 1.	Comparison of the effects of xanomeline treatment on the specific binding of [ <sup>3</sup> H]QNB and [ <sup>3</sup> H]NMS in CHO cells expressing hM3 receptors.	210
Figure 2.	Comparison of the effects of xanomeline treatment on the specific binding of [ <sup>3</sup> H]QNB and [ <sup>3</sup> H]NMS in CHO cells expressing hM1 receptors.	212
Figure 3.	Effects of blocking receptor activation on the ability of xanomeline to initiate long-term effects on [ <sup>3</sup> H]NMS saturation binding parameters in CHO cells expressing hM3 receptors.	214
Figure 4.	Time course of switching of the pharmacological profile of xanomeline from an agonist to an antagonist in CHO cells expressing hM1 or hM3 receptors.	215
Figure 5.	Effects of decreasing receptor expression using PrBCM on saturation binding parameters in CHO cells expressing hM4 receptors.	217
Figure 6.	Effects of blocking the G <sub>i/o</sub> G proteins on the ability of xanomeline to induce long-term changes in CHO cells expressing the hM4 receptor.	218

## **Appendix A: Differences observed between suspension and monolayer protocols**

Figure 1. Comparison between [<sup>3</sup>H]NMS saturation binding paradigms on assays conducted in suspension versus in monolayer in CHO cells stably expressing hM3 receptors. 246

Figure 2. Comparison between inhibition of [<sup>3</sup>H]NMS binding following xanomeline pretreatments for assays conducted in suspension versus in monolayer in CHO cells stably expressing hM1 receptors. 247

## **CHAPTER 1: Introduction**



## **Overview**

Muscarinic acetylcholine receptors are known to be involved in a variety of diseases such as Alzheimer's disease, schizophrenia, epilepsy, Parkinson's disease and depression. However, treatment of these diseases with cholinergic agents has had limited success due to side effects as a result of their non-selective binding to multiple subtypes of muscarinic acetylcholine receptors. One of the most difficult hurdles in developing agonists or antagonists that bind to muscarinic receptors with subtype selectivity is due to the highly conserved nature of the orthosteric (primary) binding domain. New research is beginning to focus on allosteric (secondary) binding sites as drug targets because these sites are less conserved across subtypes, suggesting they may lead to discovery or designing of drugs with greater subtype selectivity. Continued research is needed to learn more about these allosteric sites and the interaction of potential drugs with these sites.

Xanomeline is a novel muscarinic receptor agonist that has been shown to bind at both the orthosteric site and at an allosteric site. The major goal of this research is to characterize the actions of prolonged exposure of xanomeline at the various subtypes of muscarinic receptors. It is important to compare the long-term actions of xanomeline in the same type of cells that express individual subtypes of muscarinic receptors. This avoids complications caused by factors not related to the type of muscarinic receptor expressed, e.g. type and level of G proteins. It is also important to compare the ability of long-term treatment with xanomeline to desensitize and down-regulate various subtypes of muscarinic receptors. Differential rates or extents of receptor desensitization could contribute to agonist selectivity under chronic treatment conditions. Thus, experiments have been designed to test the hypotheses that a) xanomeline binds to all receptor subtypes in a reversible and wash-resistant manner; b) there is subtype selectivity in the

long-term effects of wash-resistant xanomeline binding; and c) xanomeline wash-resistant binding results in long-term effects similar to those seen during prolonged incubation with conventional agonists, such as internalization and down-regulation.

This introduction will address various topics that are relevant to muscarinic receptor biology and the novel muscarinic receptor agonist xanomeline, including structure, localization and function of the muscarinic receptor; the consequences of prolonged exposure to agonists; the concept of allosteric modulation of receptors; and the past and present potential for xanomeline as a treatment for Alzheimer's disease and schizophrenia.

This thesis will address the proposed hypotheses by examining the acute and long-term effects of xanomeline binding across M1-M4 receptor subtypes, the acute and long-term effects of xanomeline on function at the M1 and M3 receptor subtypes, the role of the orthosteric binding site in the long-term effects of xanomeline at the M1 and M3 receptor subtypes, and basic information about the mechanism of the long-term effects of xanomeline.

The results will provide evidence that xanomeline binds in a reversible and wash-resistant manner at all of the muscarinic receptor subtypes tested. However, there are some distinct differences in the long-term effects of xanomeline binding. Additionally, xanomeline binding is able to stimulate a functional response both in a reversible and wash-resistant manner. Interestingly, the long-term effects of exposure to wash-resistant xanomeline binding at an allosteric site are dependent on subsequent activation of the receptor at the orthosteric site. The long-term changes that are observed appear to be a result of receptor down-regulation. Further elucidation of the mode of interaction of xanomeline with various muscarinic receptor subtypes and the underlying mechanisms may lead to a better understanding of muscarinic receptor

biology. This in turn may lead to the eventual development of both muscarinic agonists and antagonists with long duration of action and perhaps better pharmacological selectivity.

### **G protein-coupled receptors**

Muscarinic acetylcholine receptors (mAChR) are part of the large G protein-coupled receptor superfamily. The G protein-coupled receptors are one of the largest families of receptors (Wess, 2005) and contain hundreds of distinct receptors (Lefkowitz et al., 1993). Endogenous ligands for the various G proteins-coupled receptors include molecules such as biogenic amines, peptides, glycoproteins, lipids, nucleotides, ions and proteases (Gether, 2000). There are three subfamilies of G protein-coupled receptors including family A or receptors related to rhodopsin and  $\beta_2$  adrenergic receptors; family B or receptors related to glucagons; and family C or receptor related to metabotropic neurotransmitter receptors (Gether, 2000). Family A is the largest of the families and the family to which muscarinic receptors belong (Gether, 2000). The molecular models of family A G protein-coupled receptors are based on rhodopsin which was the first structure crystallized (Hulme et al., 2003).

There appears to be a conservation of properties between each of the major receptor subfamilies that are coupled to G proteins that include characteristics such as amino acid sequences, topographical organization in the plasma membrane, coupling of the receptor to effectors and regulation of kinases (Lefkowitz & Caron, 1988). Interestingly, when the domains of the receptor are expressed individually they have the ability to fold on their own and then become associated to form a functional receptor (Wess, 1996).

### **Subtypes, localization and function of muscarinic receptors**

There are five subtypes of muscarinic receptors denoted M1-M5 (Caulfield & Birdsall, 1998). Muscarinic receptors are expressed both in the brain and the periphery. In the periphery, muscarinic receptors have been shown to play a role in smooth muscle contraction, glandular secretion, and cardiac rate and force (Caulfield & Birdsall, 1998). Muscarinic receptors in the central nervous system have a wide variety of functions including playing a role in sensory, cognitive, motor, behavior, and autonomic functions (Caulfield & Birdsall, 1998; Eglen, 2005; Wess, 2004). The localization of different muscarinic receptors in the brain has been determined and knockout mice have been constructed to study the function of the individual receptor subtypes. The M1, M3 and M5 receptors are primarily expressed postsynaptically, whereas the M2 and M4 receptors are generally expressed on the presynaptic neuron (Volpicelli & Levey, 2004).

The M1 receptor is expressed in the forebrain, including the cerebral cortex, hippocampus and striatum (Wess et al., 2003). The M1 receptor is involved in cognitive processing and memory (Eglen, 2005). M1 knockout mice have an increase in locomotor behavior and some deficits in working memory (Wess et al., 2003). Based on research by Anagnostaras et al (2003), it seems that M1 receptors are not required for memory formation, but are involved in processes that require interactions between the hippocampus and cerebral cortex. This suggests that M1 receptors appear to be involved in working memory and remote reference memory (Anagnostaras et al., 2003).

The M2 receptor is expressed widely in the central nervous system and in the heart and smooth muscle (Wess, 2004; Wess et al., 2003). Autoinhibition of acetylcholine release is mediated by M2 receptors in the hippocampus and cerebral cortex (Wess et al., 2003). The M2 receptor is known to play a role in agonist-induced analgesia (Wess et al., 2003). Mice that have the M2 receptor knocked out show a

decrease in smooth muscle contraction and a decrease in body temperature (Wess et al., 2003).

M3 receptors are expressed widely throughout the brain, but at lower levels than other subtypes (Eglen, 2005; Wess et al., 2003). They are expressed at high levels in the mouse hypothalamus (Wess et al., 2003). M3 receptors are also expressed in salivary glands and peripheral organs that are innervated by parasympathetic nerves (Wess, 2004; Wess et al., 2003). The M3 receptors are known to be involved in salivation, food intake, and the gastrointestinal tract (Eglen, 2005). They have also been shown to be involved in pupillary sphincter muscle contractility (Wess, 2003). Mice in which the M3 receptor has been knocked out demonstrate decreases in body weight, food intake, serum leptin levels and serum insulin levels (Yamada et al., 2001), but an increase in energy expenditure (Gautam et al., 2006). Interestingly, the knock out mice also do not have gastrointestinal problems with absorption or motility (Gautam et al., 2006).

The M4 receptor is expressed in the forebrain (Wess, 2004), neostriatum and the ventral spinal cord (Messer, 2002). The M4 receptor is involved in autoinhibition of acetylcholine release in the striatum (Wess et al., 2003). Similar to the M2 receptor, the M4 receptor is known to be involved in agonist-induced analgesia (Wess et al., 2003). The M4 receptors have also been shown to play a role in dopamine release in the brain (van Koppen & Kaiser, 2003).

The M5 receptor is found in both neuronal and non-neuronal cells (Wess, 2004). The M5 receptor is involved in facilitating dopamine release in the striatum (Wess et al., 2003). It appears that the expression of the M5 receptors parallels, at least to some degree, dopaminergic transmission (Eglen, 2005). In addition, the M5 receptor appears to be involved in dilation of cerebral arteries and arterioles (Wess, 2004).

## **Structure and binding of muscarinic receptors**

The structure of muscarinic receptors is similar across all five subtypes. Muscarinic receptors have seven transmembrane domains. The N-terminus is located extracellularly and contains multiple glycosylation sites. Interestingly, this region is not required for expression or function of the receptor (Caulfield & Birdsall, 1998). In contrast, the C-terminus is located intracellularly. The third cytoplasmic domain of the receptor interacts with the G protein (Caulfield & Birdsall, 1998; Volpicelli & Levey, 2004). Muscarinic receptors are thought to have over ninety percent conservation in the transmembrane domains between subtypes (Felder, 1995). There is a higher degree of similarity compared to the loops or the C- or N-terminus (Dohlman et al., 1991). There is also a high degree of homology between the orthosteric (primary) binding region across muscarinic receptor subtypes (Eglen, 2005).

A variety of molecules have been shown to bind to all five muscarinic receptors in a non-subtype selective manner. These include the endogenous ligand acetylcholine and other compounds including, but not limited to, muscarine, carbachol, pilocarpine, oxotremorine, scopolamine, and atropine. For example, the affinity of [<sup>3</sup>H]N-methylscopolamine was found to be similar across muscarinic receptor subtypes expressed in CHO cells (Christopoulos et al., 1999), demonstrating binding at the orthosteric site in a non-subtype selective manner.

The residues that appear to be important in the binding of positively charged ligands, such as acetylcholine at the primary binding site, are aspartate 147, tyrosine 148, tyrosine 506, tyrosine 529 and tyrosine 533 (Han et al., 2005; Wess, 1996). Another study suggests that aspartate 113 in transmembrane domain 3 is important in the binding of acetylcholine (Gether, 2000). Additional amino acids are also thought to

play a role in ligand binding (Wess, 1996). However, it is also possible that different agonists may not share an identical binding site on the receptor (Gether, 2000).

Ligand binding is thought to result in conformational changes in the receptor. Activation of muscarinic receptors is thought to occur through an interplay between transmembrane domains 3, 5, 6 and 7 (Eglen, 2005; Wess, 1996). Research was conducted to determine the specific conformation changes of the receptor when an agonist binds and activates it. Based on a study using cross-linking of cysteine residues to determine the proximity of the external portions of the transmembrane domains to each other, it was determined that the proximity of transmembrane domains 3 and 7 increases when the receptor is activated with carbachol (Han et al., 2005). Another study using cross-linking of cysteine residues found that the cytoplasmic transmembrane domains 5 and 7 move closer together upon agonist activation (Ward et al., 2002). Similar results were seen with the  $\beta$ -2 adrenergic receptor, suggesting that the mechanism of receptor activation may be conserved across class A G protein-coupled receptors (Ward et al., 2002).

Thus far, orthosteric muscarinic agonists generally have been found to have a low degree of selectivity between subtypes (Wess, 2004). Some muscarinic antagonists, however, have been shown to exhibit significant selectivity among muscarinic receptor subtypes. The commonly larger size of antagonist molecules provides accessory binding to additional receptor domains that vary in their sequence from one subtype to another. For example, pirenzepine is a muscarinic antagonist that is thought to have some selectivity for the M1 receptor compared to other muscarinic receptor subtypes (Wess, 1996).

In addition to the orthosteric binding site, all subtypes are thought to have allosteric binding sites which may be less conserved between subtypes (Eglen, 2005).

The allosteric site is spatially distinct from the site where the endogenous ligand binds (Birdsall & Lazareno, 2005). The potential for a compound that acts with more subtype selectivity through an allosteric site will be discussed further below.

### **A model of receptor activation**

The binding of an agonist to the receptor can be theoretically modeled. One type of model, the two-state model also referred to as the ternary model of binding, posits that the receptor exists in two conformations; an inactive conformation and an active conformation (Gether, 2000). In this model, the agonists are able to bind to receptors in the active conformation and thus shift the equilibrium toward that of activated receptors. Antagonists bind to both the inactive and active conformations, resulting in no change in equilibrium. In contrast, inverse agonists bind to the inactive form of the receptor resulting in a shift in equilibrium towards inactive receptors (Gether, 2000). This is quantified by the cooperativity factor  $\alpha$ , which is based on the affinity of the ligand for the active state of the receptor compared to the affinity for the inactive state (Christopoulos & Kenakin, 2002; Hulme et al., 2003). This model is taken further in the extended ternary complex model to incorporate binding of G proteins to the receptor (Hulme et al., 2003). In this case, the receptor again exists in inactive and active conformations. However, the receptor in the active state (either with bound or no ligand) now binds to the G protein (Hulme et al., 2003). It should be noted that the G protein could also bind to the inactive form of the receptor (Christopoulos & Kenakin, 2002). In this model it is also possible that multiple G proteins have the ability to bind to the active receptor conformation, which would allow for potential specificity in receptor signaling (Kenakin, 2003).



### **Receptor trafficking**

There may be some specificity in the way that different agonists activate the receptors resulting in activation of specific pathways even when the receptors are coupled to the same G proteins (Eglen, 2005). In the literature this has been referred to as “receptor trafficking”, “stimulus trafficking”, “functional selectivity”, “biased agonist” and “ligand induced differential signaling” (Kenakin, 2003; Urban et al., 2007). In this model, the receptor would exist in various active conformations and the binding of an agonist would have a specific effect on the conformation of the receptor, which ultimately could result in the activation of different G proteins or signaling cascades (Kenakin, 2003; Urban et al., 2007). If this occurs, it is likely that different agonists interact with the receptor in a way that breaks interactions between domains of the receptor in a unique manner upon activation (Urban et al., 2007). Support for this type of receptor-signaling interaction comes from the observation that different agonists have different rank-orders of intrinsic efficacy for the activation of multiple G proteins (Kenakin, 2003). Additionally, the presence of multiple active states of the receptor is supported by an agonist acting as a partial agonist in one setting and an inverse agonist in another setting when the same receptor is expressed (Kenakin, 2003; Urban et al., 2007). The difference in the ability of agonists to induce receptor internalization further supports this concept (Urban et al., 2007). Targeting activation of the receptor through specific conformations could be a potential strategy for drug development for the treatment of various diseases (Kenakin, 2003).

### **Drug-receptor interactions**

Molecules bind to the receptor in a specific manner to activate (agonists), block (antagonists) or decrease (inverse agonists) receptor activation. Agonists have both

affinity and intrinsic efficacy for the receptor, whereas antagonists have only affinity (Christopoulos & El-Fakahany, 1999). An inverse agonist has negative efficacy (Christopoulos & El-Fakahany, 1999). The action of an agonist is dependent on the system that it is being studied, thus it can only be defined as a partial or full agonist for the system in which the response is measured (Christopoulos & El-Fakahany, 1999). When a system has spare receptors this means that only a fraction of the receptors needs to be activated to get a maximal response. Additionally, it is important to note that a receptor saturating concentration of a drug in one system may not be saturating in a different system (Christopoulos & El-Fakahany, 1999).

### **Signaling pathways for muscarinic receptors**

Functionally, muscarinic receptors have been shown to activate various signaling pathways including phospholipase C, inhibition of adenylate cyclase, modulation of ion channels, activation of mitogen-activated protein kinase (involved in cell survival, differentiation and synaptic plasticity) and small GTPases (Volpicelli & Levey, 2004). Specifically, the M<sub>1</sub>, M<sub>3</sub> and M<sub>5</sub> receptors are coupled to G<sub>q</sub> and G<sub>11</sub> G proteins that leads to the activation of phospholipase C (Volpicelli & Levey, 2004). In addition M<sub>1</sub>, M<sub>3</sub> and M<sub>5</sub> receptors have also been shown to activate phospholipase A<sub>2</sub>, phospholipase D and tyrosine kinases (Wess, 1996). In comparison, M<sub>2</sub> and M<sub>4</sub> receptors are coupled to G<sub>i</sub> and G<sub>o</sub> G proteins that leads to the inhibition of adenylate cyclase (Eglen, 2005). M<sub>2</sub> and M<sub>4</sub> receptors have also been shown to increase ATR receptor-induced phosphorylation of phospholipase A<sub>2</sub> (Wess, 1996), activate inward rectifying potassium channels and inhibit activation of fast voltage sensitive calcium channels (Hulme et al., 2003). All subtypes of muscarinic receptors have been shown to activate extracellular signal related kinase (ERK) (Hulme et al., 2003). Additionally, muscarinic receptors

have been shown to couple to G proteins other than the ones to which they preferentially couple. This may occur both in endogenous systems as well as in recombinant expression systems (Nahorski et al., 1997). Furthermore, the coupling to the various G proteins may be dependent on the agonist/antagonist bound (Nahorski et al., 1997). In general, when the receptor is activated, GDP is exchanged for GTP on the G protein resulting in its dissociation from the receptor (Wess, 1996). Subsequently, the G protein is dissociated into  $\alpha$  and  $\beta\gamma$  subunits, which both activate downstream signaling cascades (Wess, 1996).

One pathway that is stimulated by receptor activation, primarily at the M1, M3 and M5 receptor subtypes, results in the production of inositol phosphates. In this pathway the  $\alpha$  and/or  $\beta\gamma$  subunits of the G protein activates phospholipase C (PLC) (Caulfield, 1993; Lanzafame et al., 2003). Interestingly, there are thought to be at least nine isoforms of PLC, but the role of agonists in stimulating the various isoforms remains unclear (Caulfield, 1993). PLC then mediates the conversion of the membrane bound phosphatidylinositol 4,5-bisphosphate ( $PIP_2$ ) into diacylglycerol (DAG) and inositol 1,4,5-triphosphate ( $IP_3$ ) (Caulfield, 1993).  $IP_3$  stimulates the release of calcium from intracellular stores within the endoplasmic reticulum (Caulfield, 1993; Felder, 1995). In one type of functional assay, the pool of  $PIP_2$  is labeled with radiolabeled myo-inositol. Following stimulation with an agonist, different metabolites of myo-inositol are formed such as glycerophosphoinositol; myo-inositol 1-phosphate; 1, 4-bisphosphate; and inositol 1, 4, 5-triphosphate which can be quantitated (Berridge et al., 1983). The levels of each of the metabolites produced depend on the time period of exposure to the agonist (Berridge et al., 1983). Subsequently, these products are converted back into  $PIP_2$  by phosphomonoesterases (Berridge et al., 1983). In parallel, DAG stimulates protein kinase C (PKC), although the levels of this stimulation appear to be minimal

when muscarinic receptor activation occurs (Caulfield, 1993). Stimulation of the effectors IP<sub>3</sub> and DAG result in alterations in cell transformation, differentiation and growth (Lanzafame et al., 2003).

Activation of phospholipase A<sub>2</sub> (PLA<sub>2</sub>) as a result of inositol phosphate-induced calcium signaling through stimulation of the M1, M3 and M5 receptors has been shown to occur (Lanzafame et al., 2003). It is also thought that stimulation of the M2 and M4 receptors can result in indirect activation of PLA<sub>2</sub> (Lanzafame et al., 2003; Wess, 1996). PLA<sub>2</sub> stimulation results in the production of arachidonic acid and lysophospholipid as a result of phospholipid hydrolysis (Lanzafame et al., 2003). PLA<sub>2</sub> has been shown to play a role in membrane fluidity, and thus changes in PLA<sub>2</sub> activity may play a role in diseases such as Alzheimer's disease (Lanzafame et al., 2003).

Adenylate cyclase (AC) activity is modulated as a result of receptor stimulation by agonists. There have been shown to be at least ten isoforms of AC (Lanzafame et al., 2003), and the result of receptor activation may be dependent on which isoform of AC is present (Lanzafame et al., 2003). Generally, agonist binding to the M2 or M4 receptors results in inhibition of AC through the  $\alpha$  subunit of the G<sub>i/o</sub> G proteins (Caulfield, 1993; Lanzafame et al., 2003). Subsequently, this results in a decrease in levels of cAMP (Caulfield, 1993). Furthermore, the decrease in cAMP results in a decrease in protein kinase A (PKA) activity. In contrast, it has been demonstrated that the M1, M3 and M5 receptors can also stimulate minimal levels of AC, resulting in an increase in cAMP levels and an increase in PKA activity (Caulfield, 1993; Felder, 1995). It is possible that the accumulation of cAMP following receptor activation could be a result of calcium or protein kinase C signaling (Felder, 1995; Nathanson, 2000).

Mitogen-activated protein kinase (MAPK) is divided into multiple pathways, one of which is extracellular signal regulated kinase (ERK) (Lanzafame et al., 2003). ERK

has been shown to influence cell growth, differentiation and survival (Werry et al., 2005). ERK signaling has been shown to greatly depend on protein-protein interactions. These interactions are important in allowing the signaling pathway to be complex; controlling the order and orientation of substrates and enzymes; and targeting the signaling to specific subcellular locations (Werry et al., 2005). ERK activation is thought to occur as a result of agonist stimulation of all subtypes of muscarinic receptors (Lanzafame et al., 2003). When expressed in COS-7 cells both M1 and M2 receptors have been shown to stimulate ERK following exposure to carbachol (Crespo et al., 1994). In further support of the idea of the activation of ERK by agonists, stimulation of the M2 receptor has been shown to activate both Raf and Ras, which are upstream of ERK (Winitz et al., 1993). Thus, this activation occurs for both  $G_q$  and  $G_i$  G proteins. The activation of ERK through the M1, M3 and M5 muscarinic receptors is thought to be a result of PKC activation that is downstream of PLC activation (Lanzafame et al., 2003). In the case of M2 and M4 muscarinic receptors, activation of ERK appears to be dependent on the  $\beta$  subunit of the G protein (Lanzafame et al., 2003). The MAPK maximal response to agonist stimulation was observed between five and ten minutes; however, the response remained elevated for up to 90 minutes after agonist stimulation (Wotta et al., 1998). The activation of MAPK subsequent to stimulation of the receptor with an agonist is time and concentration dependent (Wotta et al., 1998).

There are thought to be six different G protein-coupled receptor kinases (GRKs) (Pitcher et al., 1998) that are involved in the desensitization of the response to agonist stimulation (Wess, 1996). They only phosphorylate G protein-coupled receptors when the receptors are activated by the binding of an agonist, which may have to do with the conformation of the receptor (Pitcher et al., 1998; Wess, 1996). Phosphorylation of the muscarinic receptors by GRKs is thought to occur on serine and threonine residues on

the third cytoplasmic loop of the receptor and possibly on the C-terminus (van Koppen & Kaiser, 2003). Phosphorylation by GRKs ultimately may result in sequestration of the receptor through the binding of arrestins (van Koppen & Kaiser, 2003; Wess, 1996). The phosphorylation by GRKs is thought to occur within minutes of the receptor being activated by an agonist (Pitcher et al., 1998). However, sequestration as a result of GRK phosphorylation does not appear to occur at all types of G protein-coupled receptors (Wess, 1996).

### **Consequences of prolonged exposure of muscarinic receptors to agonists and antagonists**

Prolonged exposure of muscarinic receptors to agonists can result in receptor desensitization through mechanisms such as internalization, down-regulation, uncoupling of the G protein from the receptor and structural changes such as phosphorylation of the receptor (Maloteaux & Hermans, 1994). Initially receptor desensitization may occur without receptor internalization (Maloteaux & Hermans, 1994). When receptor internalization does occur cytoskeletal elements are thought to be important both for internalization and reinsertion of receptors into the cellular membrane (Maloteaux & Hermans, 1994). Acute desensitization results in a rapid decrease in the response with the return of the response occurring without new receptor synthesis (Dohlman et al., 1991). In contrast, long-term desensitization results in a decrease in receptor expression over time that requires protein synthesis to replace the deficit (Dohlman et al., 1991). This suggests that the decrease in muscarinic receptor expression occurs in two phases. The first is internalization in which the receptors can still be bound to the G protein and can subsequently be reinserted into the membrane. This is followed by down-regulation of the receptor (El-Fakahany & Cioffi, 1990).

Receptor expression can be measured with the aid of radiolabeled receptor antagonists. Two of these antagonists are [<sup>3</sup>H]N-Methylscopolamine ([<sup>3</sup>H]NMS) and [<sup>3</sup>H]quinuclidinyl benzilate ([<sup>3</sup>H]QNB). NMS is a permanently charged amine which can only bind to receptors expressed on the cell surface (Feigenbaum & El-Fakahany, 1985), whereas QNB is a lipophilic antagonist which can bind to receptors on the cell surface or internalized receptors that have not yet been reinserted or degraded (Ehlert et al., 1981; Vickroy & Malphurs, 1994).

When cells are treated with carbachol for prolonged periods of time there is a time dependent decrease in the levels of cell-surface receptor expression determined by both [<sup>3</sup>H]NMS and [<sup>3</sup>H]QNB binding; however, there was no change in the affinity of either radioligand (Cioffi & El-Fakahany, 1989; Hu et al., 1991). In contrast, brief treatment of cells expressing the M2 receptor with carbachol for 30 minutes resulted in a decrease in [<sup>3</sup>H]NMS binding, but no change in [<sup>3</sup>H]QNB binding, suggesting that the receptors are being internalized (Tsuga et al., 1998). This suggests that in binding studies, [<sup>3</sup>H]NMS is more beneficial in detecting rapid changes in receptor expression on the cell surface compared to [<sup>3</sup>H]QNB (Feigenbaum & El-Fakahany, 1984). When cells expressing M2 receptors were treated with carbachol, the maximal decrease in receptor expression resulted after 3 hours of treatment (May et al., 2005). In addition, Stope et al. (2003) demonstrated that treatment of cells expressing the M3 receptor with carbachol for 24 hours resulted in an approximately 50 percent reduction in the expression levels of the G<sub>q/11</sub> G proteins. When cells expressing the M1 receptor were exposed to carbachol, the G<sub>q</sub> expression decreased to 50 percent of the maximum occurred even more rapidly (after 3 hours); however, in this case the treatment was found to have no effect on other G proteins (Mullaney & Milligan, 1993). A decrease in G<sub>q/11</sub> G protein expression independent of receptor down-regulation was also observed

by van de Westerlo et al (1995) in response to prolonged agonist treatment. Furthermore, work by Shifrin & Klein (1980) demonstrated that prolonged continuous activation or intermittent activation over time with carbachol results in a similar decrease in receptor expression on the cell surface. When atropine was used to block the orthosteric site during prolonged exposure to agonist there no longer was a decrease in receptor expression, demonstrating that the decrease in receptor expression was the result of agonist binding to the receptor (El-Fakahany & Cioffi, 1990). In contrast, the effect of prolonged pilocarpine, another muscarinic receptor agonist, treatment on receptor expression is less clear. Milligan & Strange (1983) found that there was no change in receptor expression following prolonged treatment with pilocarpine (6 hours) as measured by [<sup>3</sup>H]NMS binding.

Agonist binding to the receptor can also influence the production of mRNA encoding the receptor. When CHO cells expressing M1 receptors were exposed to agonists, there was a reduction in mRNA that was associated with the agonist-induced down-regulation of the receptors (Wang et al., 1990). Additionally, the decrease in mRNA expression was shown to require a functional response because blocking the receptor with atropine during carbachol treatment resulted in only a minimal decrease in mRNA levels (Wang et al., 1990). Similarly, when cells expressing M3 receptors were exposed to carbachol for a prolonged period of time, there was a decrease in the expression levels of mRNA; however, the decrease occurred well after the decrease in cell-surface receptor expression (Fukamauchi et al., 1993; Steel & Buckley, 1993). Thus, the decrease in receptor expression levels following agonist treatment is most likely due to a combination of increased receptor degradation and a decrease in receptor production as a result of the decrease in mRNA expression (Wang et al., 1990).



There appears to be a correlation between agonist efficacy and the ability of the agonist to cause desensitization of the response (Hu et al., 1991). Furthermore, the desensitization of phosphoinositide (PI) hydrolysis appears to parallel the decrease in receptor expression (Thompson & Fisher, 1990). However, the time course of this response may depend on the conditions used. Coiffi & El-Fakahany (1989) found that the decrease in the ability of the agonist to stimulate inositol phosphate production occurred more slowly than down-regulation of cell-surface receptor density. For example, prolonged treatment of cells with the full agonist carbachol resulted in a time-dependent decrease in PI hydrolysis subsequent to further stimulation with carbachol (Hu et al., 1991; Mullaney & Milligan, 1993). Additionally, no increase in basal levels of PI hydrolysis was observed following carbachol treatment (Cioffi & El-Fakahany, 1989). In contrast, when cells were treated with carbachol followed by subsequent stimulation with the partial agonist McN-A-343, the decrease in response was more pronounced than that observed with carbachol (Hu et al., 1991). In addition, when McN-A-343 was used in place of carbachol in the pretreatment paradigm it was still able to desensitize its own response, but to a lesser degree than was observed with carbachol (Hu et al., 1991).

Research by van de Westerlo et al. (2005) was conducted to determine what regions of the receptor are important for receptor down-regulation in response to prolonged agonist stimulation. They demonstrated that when CHO cells expressing the M3 receptor were exposed to carbachol for prolonged periods of time the receptors were down-regulated; however, if a threonine to alanine mutation was made in the C-terminus of the receptor, down-regulation of the receptor no longer occurred. Additionally, although the threonine residues located at positions 550, 553 and 554 in the C-terminus appear to be important in receptor down-regulation they do not appear to play a role in

receptor binding or stimulation of signaling cascades (Yang et al., 1993). It is possible that these residues are sites for PKC phosphorylation; however, using an inhibitor to block phosphorylation in wild-type cells does not prevent receptor down-regulation (Yang et al., 1993).

Antagonists are not generally thought to induce receptor internalization (Maloteaux & Hermans, 1994). In contrast to treatment with agonists, long-term administration of atropine, a known muscarinic receptor antagonist, in rats (injections for seven days) resulted in an increase in [<sup>3</sup>H]QNB binding in the cortex, hippocampus and heart (Lee & Wolfe, 1989). In addition, in a cell culture system, treatment of cells expressing the M2 receptor with antagonists, atropine (20 nM) or gallamine (20 μM), resulted in an increase in cell-surface receptor density with no change in radioligand affinity (May et al., 2005). Similarly, when cells expressing the M3 receptor were used, there was an increase in receptor expression following 24 hour exposure to atropine followed by washing (Ward et al., 2002). However, the increase in receptor expression occurred more slowly (over 12 hours) compared to the time needed for a decrease in receptor expression subsequent to prolonged treatment with an agonist such as carbachol (May et al., 2005). It should be noted that atropine was found to be an inverse agonist, rather than an antagonist, when studied using a constitutively active M3 receptor; thus, the effects on receptor density may be slightly different depending on the system (Thor et al., 2008). When prolonged *in vivo* treatment (injections for seven days) with another muscarinic receptor antagonist, scopolamine, was tested, it also resulted in an increase in cell-surface receptor density with no change in radioligand binding as measured by [<sup>3</sup>H]QNB binding in tissues obtained from rat brain (Ben-Barak & Dudai, 1980). Together these results demonstrate that antagonist treatment can result in

upregulation of cell-surface receptor density, as opposed to agonist treatment down-regulating muscarinic cell-surface receptor density.

### **Influence of receptor mutations and expression levels on receptor activity**

It should be noted that some G protein-coupled receptors have been shown to have endogenous activity in the absence of agonist stimulation. This activity can be a result of receptor mutations or changes in the expression level of the receptors. For example, some receptor mutations have been shown to result in high levels of constitutive receptor activity (Gether, 2000; Lefkowitz et al., 1993). Research has also demonstrated that mutations in the receptor can change the function of an antagonist to an agonist (Thor et al., 2008). Interestingly, although the amino acid sequence of muscarinic receptors appears to be conserved across species, minor differences in the amino acid sequence can result in dramatic changes in the pharmacological profile observed between different species (Caulfield & Birdsall, 1998).

Additionally, in *in vitro* receptor expression systems, high receptor expression can lead to elevated levels of basal receptor activation (Christopoulos & Kenakin, 2002; Milligan et al., 1995), possibly because more receptors will spontaneously be in the active state of the receptor (Chen et al., 2000). It has also been shown that partial agonists produce larger responses when there is high receptor expression or the receptor is more efficiently coupled to signaling cascades (Hu & El-Fakahany, 1990). Interestingly, various compounds have been shown to act as a partial agonist under one set of conditions but a full agonist under other conditions (Milligan et al., 1995). Thus, differences in responses can be observed for a given agonist depending on the level of receptor expression and the response system (Wang & El-Fakahany, 1993).

### **In vitro receptor expression systems**

The use of receptor expression systems to study the properties of receptor binding and function can be beneficial. Some of the benefits of employing a cell culture system are that the cells are homogeneous, receptors can be expressed individually, cells are cultured in monolayer making it easier to administer drugs, it is possible to perform simultaneous experiments on receptor density and function and in intact cells, cellular localization of the receptor can be studied (El-Fakahany & Cioffi, 1990). The use of receptor expression systems is important because various tissues express multiple receptor subtypes, making it difficult to determine the properties of individual receptor subtypes (Bonner, 1989).

However, there are also potential problems with recombinant receptor expression systems. One problem is that in a recombinant expression system the receptor or the G protein may be interacting with membrane or cytosolic proteins that they do not endogenously associate with (Christopoulos & El-Fakahany, 1999; Kenakin, 1997). It is also possible that receptor overexpression in an experimental paradigm could lead to a compound being classified as a partial agonist in the experimental system, but when it is part a complex system it functions more like an antagonist (Christopoulos & El-Fakahany, 1999). Another potential problem is that the G protein levels may be low compared to the receptor expression levels, thus resulting in a lack of change of the receptor to a high affinity state which facilitates binding (Kenakin, 1997). It is possible that binding studies may be better determinants than functional studies because the result of binding studies will be an increased signal whereas in functional studies the maximal response can be saturated (Kenakin, 1997). Lastly, it may be possible to have selectivity *in vivo*, which is not observed *in vitro*, due to differences in the tissue such as affinity, efficacy, receptor number, and receptor-response coupling (Wood et al., 1999).

When using a receptor expression system it is also important to determine if the receptors of interest are expressed endogenously in the system. One cell line that is commonly used as an expression system for muscarinic receptors is Chinese hamster ovary (CHO) cells. When wild-type CHO cells were transfected with nitric oxide synthase there was a response to carbachol that was blocked by atropine, suggesting the presence of endogenous muscarinic receptors (Wang et al., 1995). However, muscarinic receptors were not detected with radioligand binding assays or functional assays looking at inositol phosphate production (Wang et al., 1995). Thus, this cell line is used to study muscarinic receptor biology, with the assumption that the endogenous expression of muscarinic receptors is negligible compared to the expression levels of the transfected muscarinic receptors.

Additionally, the experimental conditions of the assay need to be considered when using receptor expression systems, since differences have been observed between assays conducted with cells that are attached to a surface (monolayer) versus floating in buffer (suspension). For example, when forskolin-stimulated cAMP accumulation was studied there was a difference in the response when the assay was conducted in suspension versus monolayer. The assay conducted in suspension only resulted in 75 percent of the response obtained when cells were assayed in monolayer. In contrast, the response to pilocarpine was larger in cells in suspension compared to those assayed in monolayer (Mistry et al., 2005). This suggests that the response is dependent both on the agonist and the assay conditions. Additionally, when cells were treated with agonists and the level of receptor expression studied in monolayer or suspension there was a greater loss of cell-surface receptors for cells treated in monolayer compared to those in suspension as measured by [<sup>3</sup>H]NMS binding (Thompson & Fisher, 1990). The difference between assays and the possible causes

for these observed differences are discussed further in Appendix A. In this thesis assays will be conducted in monolayer unless otherwise noted.

### **Allosteric interactions with the receptor**

Allosteric sites have been found on a variety of receptors including ligand-gated ion channels, nicotinic receptors, GABA<sub>A</sub> receptors and G protein-coupled receptors (Gao & Jacobson, 2006). There are different types of allosteric modulators. Allosteric enhancers result in a conformation change at the orthosteric site that results in higher affinity for the ligand; allosteric agonists result in the G protein coupling more easily to the receptor irrespective of the action on the orthosteric site; and allosteric antagonists result in a decrease in receptor affinity and/or make the G protein less likely to couple to the receptor (Christopoulos & Kenakin, 2002; Wess, 2005).

Compounds that bind in an allosteric manner can result in conformational changes in the receptor which can result in the modulation of receptor activity (Christopoulos, 2002). In allosteric interactions, the binding of one ligand affects the ability of another ligand to bind. This occurs in a reciprocal manner between the two ligands (Birdsall et al., 1996). Changes in receptor conformation can result in changes in receptor affinity. Positive cooperativity results in an increase in affinity, negative cooperativity results in a decrease in affinity and neutral cooperativity results in no change in affinity (Christopoulos, 2002). The action of a particular allosteric modulator may be ligand specific. For example, WIN51708 displayed positive cooperativity with [<sup>3</sup>H]NMS binding, however it had negative cooperativity with [<sup>3</sup>H]QNB binding (Lanzafame et al., 2006). Thus, it may be possible to get some degree of subtype selectivity based on cooperativity rather than affinity (Lanzafame et al., 2006). In addition, it is important to consider that the effect on receptor function as a result of

allosteric modulation is not always the same as it is on binding (Christopoulos, 2002). It is also important to consider that the binding of allosteric ligands to the receptor can be dependent on the experimental conditions such as ionic strength, divalent cations and temperature (Birdsall & Lazareno, 2005).

Interaction of a compound through an allosteric site may be more advantageous than acting at the orthosteric site because it may be possible to get greater subtype selectivity since allosteric sites likely have greater sequence divergence across receptor subtypes. If the effect of the allosteric modulator is saturable it could prevent problems with overstimulation. Additionally, the activation of the response could be manipulated so the response occurs only when the endogenous agonist is present resulting in tissue specific activation (Christopoulos, 2002; Christopoulos & Kenakin, 2002).

### **Pharmacological profile of allosteric binding**

Allosteric modulation of receptors results in changes in the binding and functional properties determined using various receptor assays. One classic hallmark of allosteric modulation is the inability to decrease binding to non-specific levels when employing an inhibition of radioligand binding assay. This is dependent on the concentration of the radioligand (Christopoulos & Kenakin, 2002; Lee & El-Fakahany, 1991). Additionally, using a saturation binding paradigm, the shift in binding is saturable with increasing concentrations of the allosteric agent (Christopoulos & Kenakin, 2002). Allosteric binding also has the ability to change the association and/or dissociation kinetics of a reaction at the orthosteric site, which can change the time that it takes the system to reach equilibrium (Birdsall & Lazareno, 2005; Christopoulos, 2002; Christopoulos & Kenakin, 2002). In radioligand binding assays positive cooperativity is determined using

a radioligand concentration below the  $K_d$ , whereas negative cooperativity is determined using a high concentration of radioligand (Birdsall & Lazareno, 2005).

### **Allosteric sites on muscarinic receptors**

It is possible that there are multiple allosteric binding sites on muscarinic receptors (Christopoulos & Kenakin, 2002; Wess, 2005). It is also possible that a single receptor can have multiple allosteric modulators bound at different sites at the same time (Christopoulos & Kenakin, 2002). It has been suggested that a possible allosteric site on muscarinic receptors is located near the orthosteric site, but more extracellularly (Christopoulos & Kenakin, 2002). It is also likely that muscarinic receptors have specific allosteric sites because some allosteric modulators are able to compete with each other (Christopoulos & Kenakin, 2002). These allosteric sites may overlap across receptor subtypes or have varying degrees of subtype selectivity.

A variety of compounds have been found to be allosteric modulators of muscarinic receptors. These include compounds such as neuromuscular blockers, reactivators of cholinesterases, ion channel blockers, and local anaesthetics, among others (Tucek & Proska, 1995). One such compound is gallamine. Gallamine is thought to be an allosteric modulator at all subtypes of muscarinic receptors (Clark & Mitchelson, 1976; Wess, 1996). However, the effects appear to be the most pronounced at the M2 receptor subtype (Wess, 1996). It was able to inhibit responses to both acetylcholine and carbachol in the guinea pig atria (Clark & Mitchelson, 1976). Gallamine decreased both the association and dissociation of additional ligands at the orthosteric binding site (Tucek & Proska, 1995; Wess, 1996). Gallamine was not able to completely inhibit [ $^3$ H]NMS binding, which is a hallmark of an allosteric modulator (Spalding et al., 2006). In a saturation binding paradigm, gallamine reduced [ $^3$ H]NMS binding affinity in a



manner that was distinct from a conventional reversible agonist (Lee & El-Fakahany, 1991). Additionally, gallamine may mediate changes in the receptor conformation resulting in positive cooperativity with the G protein, but negative cooperativity with the agonist binding site (Christopoulos & Kenakin, 2002). It is thought that gallamine binds at an extracellular location on the receptor, perhaps extracellularly to the agonist binding site in a manner in which it is able to at least partially block the orthosteric site (Birdsall et al., 1996).

Alcuronium is another allosteric modulator of muscarinic receptors. It has been shown to inhibit the dissociation of [<sup>3</sup>H]QNB from the receptor and inhibit both the association and dissociation of [<sup>3</sup>H]NMS from the receptor (Tucek & Proska, 1995). However, its effects on binding of [<sup>3</sup>H]NMS are dependent on the tissues being studied. Alcuronium resulted in an increase in radioligand affinity in rat heart, ileum and cerebellum, which are tissues that are thought to express primarily M<sub>2</sub> and M<sub>4</sub> receptors, but a decrease in affinity in the cerebral cortex and salivary glands which are thought to express primarily M<sub>1</sub>, M<sub>3</sub> and M<sub>5</sub> receptors (Tucek & Proska, 1995). Similar to gallamine, alcuronium may bind to a region on the receptor that blocks further access to the orthosteric site either directly or through steric modifications (Tucek & Proska, 1995). One possible explanation for the discrepancy observed between the binding kinetics versus the affinity of alcuronium is that allosteric binding in this case results in changes in the affinity of the receptor that may be due specifically to conformational changes, while a change in kinetics may be the result of steric hindrance (Tucek & Proska, 1995).

Strychnine acts as an allosteric modulator of muscarinic receptors. For example, at the M<sub>4</sub> receptor it has been shown to inhibit acetylcholine binding, but enhance [<sup>3</sup>H]NMS binding (Birdsall et al., 1996). Additionally, strychnine has been shown to

decrease [<sup>3</sup>H]NMS dissociation at the M4 receptor (Birdsall et al., 1996). Strychnine has also been shown to change the kinetics at the other receptor subtypes (Birdsall et al., 1996).

AC-42 (4-*n*-butyl-1-[4-(2-methylphenyl)-4-oxo-1-butyl]-piperidine hydrogen chloride) is an example of a novel muscarinic agonist that is thought to have functional activity at only the M1 receptor subtype (Spalding et al., 2002). However, it was able to inhibit [<sup>3</sup>H]NMS binding at all subtypes of muscarinic receptors with a low potency (Spalding et al., 2002). Interestingly, the activation of the M1 receptor by AC-42 occurs at a region other than that used by carbachol, demonstrating that it is binding in an allosteric manner (Spalding et al., 2002). Mutation data for regions in the third transmembrane domain suggest that AC-42 may bind closer to the extracellular space, whereas carbachol binds deeper in the transmembrane domain of the receptor (Spalding et al., 2006). Furthermore, the N-terminus of the receptor was shown to be necessary to obtain a functional response to AC-42 (Spalding et al., 2002).

Recent research has resulted in further development of allosteric modulators of muscarinic receptors with subtype selectivity. One such drug is 1-(1'-2-methylbenzyl)-1,4'-bipiperidin-4-yl)-1H benzo[d]imidazol-2(3H)-one (TBPB) (Jones et al., 2008). It is structurally different than any of the other allosteric modulators that are currently known. TBPB has been shown to selectively activate the M1 receptor in an allosteric manner (Jones et al., 2008). When cells were treated with increasing concentrations of atropine there was a decrease in the ability of TBPB to stimulate a response, as well as a rightward shift in the functional curve, which is an indication of allosteric modulation (Jones et al., 2008). Interestingly, TBPB has been shown to potentiate NMDA currents in the hippocampus which is thought to be mediated by M1 receptor activation (Jones et

al., 2008). However, TBPB had some antagonist activity at the D2 dopamine receptor, which complicates any potential *in vivo* application (Jones et al., 2008).

Another recent report presented data on the development of a set of compounds that are positive allosteric modulators at the M4 receptor. In functional assays these compounds had no activity on their own but potentiated the response to acetylcholine at the M4 receptor (Shirey et al., 2008). However, there was no response alone or in conjunction with additional agonist to these compounds at any of the other muscarinic receptor subtypes (Shirey et al., 2008). It appears that these compounds may increase the affinity of the orthosteric site to acetylcholine and/or through positive cooperativity with the G protein (Shirey et al., 2008).

### **Muscarinic receptors, allosteric sites and diseases**

Muscarinic acetylcholine receptors are thought to play a role in a variety of diseases of the central nervous system such as Alzheimer's disease, Parkinson's disease, depression, epilepsy and schizophrenia (Wess, 2004). The potential role of muscarinic receptors in Alzheimer's disease and schizophrenia will be discussed below. The possible use of a compound that acts as an allosteric modulator of muscarinic receptor will be discussed as a possible treatment for these diseases.

### **Alzheimer's disease**

Alzheimer's disease (AD) is a neurodegenerative disease that leads to a progressive loss of cognitive function caused by neuronal loss in the hippocampus and various neocortical regions (Keller, 2006; Kotilinek et al., 2002). It is the most common cause of dementia in the elderly with the clinical characteristics of AD involving changes in behavior and personality (Minghetti, 2005; Reddy, 2006). It has been estimated that

approximately four million people may currently have AD (Campbell, 2004), a number that is projected to increase to nearly 13.5 million by 2050 (Hodes, 2006). AD is the eighth leading cause of death in the United States (Hodes, 2006). Although the exact cause of AD is unknown, the majority of AD is thought to be a result of sporadic mutations, with only two percent of cases attributed to a familial form (Reddy, 2006).

Although currently AD can only definitively be diagnosed post-mortem, it is clinically diagnosed based on clinical and neuropsychological exams and by the exclusion of other forms of dementia (Maccioni et al., 2004). Prior to being diagnosed with AD, patients have mild cognitive impairment (MCI), during which some beta amyloid plaques and neurofibrillary tangles are present. Interestingly, not all patients that have MCI progress to AD (Ashe, 2005; Blennow, 2004). The most common symptoms of AD include a decline in memory, language, visuospatial orientation, and possibly depression (Maccioni et al., 2004). Additionally, the loss of olfaction may occur before the cognitive and behavioral decline in patients with AD (Suzuki et al., 2004).

Although the exact cause of AD remains unclear, there are multiple changes or deficits observed in postmortem analysis of the brains of Alzheimer's disease patients such as the loss of cholinergic neurons (Francis et al., 1999), accumulation of  $\beta$ -amyloid plaques (Hardy et al., 1998), formation of tangles of hyperphosphorylated tau (Hardy et al., 1998), changes in various receptor expression (muscarinic receptors, Flynn et al., 1995; nicotinic receptors, Messer, 2002) and others.

### **Cholinergic hypothesis of Alzheimer's disease**

One characteristic of AD is the loss of cholinergic neurons in the basal forebrain that project to the hippocampus (Francis et al., 1999). Acetylcholine is synthesized from acetyl-CoA and choline by the enzyme choline acetyltransferase (Raedler et al., 2007).

Acetylcholine is created in the neurons and stored in presynaptic vesicles before release (Raedler et al., 2007). In AD there is a decrease in the levels of choline acetyltransferase which appears to correlate with the degree of cognitive impairment (Francis et al., 1999). The regions that seem to be the most affected by cholinergic degeneration are pyramidal neurons, nucleus basalis of Meynert, medial septum, raphe nuclei and the locus coeruleus (Francis et al., 1999). At the end-stages of AD there may be up to a 90 percent loss of cholinergic neurons that project from the basal forebrain to cortical and hippocampal areas (Yan & Feng, 2004). Interestingly, in terms of receptor expression, the presynaptic M2 receptors are lost (Flynn et al., 1995; Francis et al., 1999); however, the postsynaptic M1 and M3 receptors appear to remain at normal levels (Eglen, 2005; Flynn et al., 1995). Additionally, the levels of presynaptic M4 receptors were increased whereas the level of postsynaptic M5 receptors remained unchanged (Flynn et al., 1995). Although the number of M1 receptors remains the same between people with AD and control subjects, the binding affinity may not be the same (Flynn et al., 1991). Interestingly, the loss of the ability of muscarinic cholinergic agonists to bind occurs in brain regions that are associated with changes in AD, but not in unassociated brain regions (Flynn et al., 1995). For example, Flynn et al. (1991) demonstrated that the ability of carbachol to bind to M1 receptors with high-affinity in the cortex was reduced in AD; however, the low affinity binding remained unchanged. This change may be a result of changes in M1 receptor-effector coupling (Flynn et al., 1991). Additionally, the potential change in function of the M1 receptors in AD may be at least part of the reason why cholinergic therapies are not always effective treatments for AD (Flynn et al., 1991).

## **Cholinergic treatments for Alzheimer's disease**

In AD the impaired cholinergic neurotransmission is thought to result in deficits in memory and cognition (Zimmermann et al., 2005). The degeneration of presynaptic cholinergic neurons results in a decrease in the levels of acetylcholine present in the synapse, which further results in a decrease in the stimulation of the postsynaptic neuron (Oda, 1999). However, the postsynaptic receptors are thought to be relatively intact (Flynn et al., 1995). The first treatment strategy that was employed was to inhibit the breakdown of acetylcholine into acetate and choline by acetylcholinesterases in order to increase levels of acetylcholine in the synapse (Mesulam, 2004). Thus, the compounds that were initially developed and used in the treatment of Alzheimer's disease were acetylcholine esterase inhibitors (AChEI's) such as tacrine (Cognex), donepezil (Aricept), rivastigmine (Exelon) and galanthamine (Reminyl) (Disterhoft & Oh, 2006). One potential problem with this type of treatment is that it relies on release of acetylcholine from presynaptic nerve terminals, which is an issue because the presynaptic terminals have been shown to degenerate (Francis et al., 1999). Additionally, the higher levels of acetylcholine in the synapse not only stimulate the M1 receptors which are the desired target but also other muscarinic receptors subtypes which are thought to mediate the side-effects of treatments such as nausea, vomiting, fainting, weight loss, muscle fatigue and bradycardia (Bianchetti et al., 2006). A recent clinical trial suggested that transdermal application of these types of drugs may result in amelioration of the cognitive and behavioral symptoms of AD without some of the side effects observed with oral administration. Currently, the reason for the decrease in side effects following transdermal drug application is unclear. However, more research is needed before this could become a wide spread treatment option (Chan (a) et al., 2008).

The second type of strategy that was employed to treat AD was the use of cholinergic agonists to replace the deficiency in endogenous acetylcholine. Such agonists would be beneficial because they do not rely on presynaptic cholinergic projections to be present (Bymaster et al., 1997). However, it is necessary to find an agonist that selectively activates the M1 receptor to avoid unwanted side effects. The creation of various M1 selective agonist compounds has been attempted. These include compounds such as AF102B, AF150, talsaclidine and xanomeline (Fisher et al., 1996; Messer, 2002). Treatment with these compounds resulted in cognitive improvements measured by the AD assessment scale; however, cholinergic side effects were seen similar to those observed with acetylcholinesterase inhibitors treatment making them impractical as clinically relevant treatments for AD (Fisher et al., 1996). The results of treatment with xanomeline will be discussed in more detail below. However, it is possible that a new therapeutic could be developed in which there is greater subtype selectivity or there is positive cooperativity between the drug and acetylcholine which would facilitate the endogenous response (Jakubik et al., 1997). A drug with positive cooperativity would allow a tissue specific response and decrease the side effects normally observed with treatment (Jakubik et al., 1997).

### **Non-cholinergic treatments for Alzheimer's disease**

AD results from a compilation of many different factors. Thus, additional treatments have been tried for AD that target potential aspects of the disease that are not directly related to cholinergic transmission. There is some evidence for increases in oxidative species or inflammation, thus, treatment strategies have included antioxidant and anti-inflammatory therapy (Zimmermann et al., 2005). Additionally, a particular form of the apolipoprotein E, the type 4 allele, is thought to play a role in AD (Corder et al.,

1993). Treatment with statins were used to try to address this problem (Zimmermann et al., 2005). Another treatment is the use of an NMDA receptor antagonist, memantine (Namenda), which inhibits calcium from entering neurons (Disterhoft & Oh, 2006). This effect may be independent or in accordance with changes in cholinergic transmission. Furthermore, the accumulation of non-soluble  $\beta$ -amyloid peptides in AD has been shown to be neurotoxic. A relatively new treatment strategy employs the use of a vaccine against  $\beta$ -amyloid with the goal being to decrease the levels of  $\beta$ -amyloid deposits in the brain (Wisniewski & Konietzko, 2008). This vaccine showed great promise in mouse models of AD. However, it had mixed results when administered to human subjects; some patients developed an immune response that decreased  $\beta$ -amyloid deposition whereas others had no apparent immune response (Wisniewski & Konietzko, 2008). Although multiple approaches for the treatment of AD may be beneficial these types of treatments based on non-cholinergic targets have not been approved for clinical use and will not be further addressed here.

### **Schizophrenia**

Schizophrenia is a mental illness that is generally characterized by behavioral symptoms such as psychosis, apathy, social withdrawal and disorganized speech (Mueser & McGurk, 2004; Raedler et al., 2007). Typically people with schizophrenia have delusions and hallucinations (Mueser & McGurk, 2004). These are thought to be due to hyperactivity of dopamine signaling in the brain (Bymaster et al., 2002). In addition, schizophrenics also have cognitive deficits in learning and memory, problem solving and problems with attention (Mueser & McGurk, 2004). There are thought to be both genetic and environmental components that lead to schizophrenia (Mueser & McGurk, 2004).



Schizophrenia can be diagnosed using different types of imaging. Schizophrenic patients have larger ventricles compared to unaffected people (Mueser & McGurk, 2004). Positron emission tomography has shown that there is a change in blood flow to the thalamus and cerebellum (Mueser & McGurk, 2004). Additionally, functional magnetic resonance imaging suggests that there are global changes in brain activity rather than changes that are specifically localized to particular brain regions (Mueser & McGurk, 2004). These changes may be a result of changes that occurred during development, resulting in abnormal connectivity in the brain (Mueser & McGurk, 2004).

It is thought that problems with dopamine signaling are involved in schizophrenia. When the D2 dopamine receptor is antagonized, there is a decrease in symptoms such as delusions and hallucinations (Raedler et al., 2007). To further support this idea, imaging studies conducted on schizophrenics have demonstrated that there is an increase in dopamine neurotransmission in their brains (Raedler et al., 2007). However, changes in dopamine transmission can not account for all of the symptoms of schizophrenia; thus, it has been proposed that other neurotransmitter systems may also play a role (Raedler et al., 2007).

Studies have shown that there is an interaction between the dopaminergic and cholinergic systems in the brain, although the specifics of the interactions are still unclear (Raedler et al., 2007). In schizophrenic patients, a reduction in the numbers of muscarinic and nicotinic acetylcholine receptors have been found in the cortex and hippocampus (Friedman, 2004). Additionally, there is some evidence that there is also a decrease in muscarinic receptors in the ventral striatum (Raedler et al., 2007). These decreases are thought to be primarily in the M1 and M4 receptor subtypes (Raedler et al., 2007).

## **Treatment for schizophrenia**

Schizophrenia is frequently treated pharmacologically with antipsychotics. However, these drugs can have negative side-effects in terms of cognition, extrapyramidal symptoms and tardive dyskinesia (Mueser & McGurk, 2004). There are two classes of drugs that are frequently used in treatment, typical and atypical antipsychotics (Bymaster et al., 2002). The most commonly used typical antipsychotics are haloperidol and chlorpromazine, which are dopamine antagonists (Bymaster et al., 2002). In contrast, the atypical antipsychotics are thought to be serotonin/dopamine antagonists and include drugs such as risperidone and clozapine (Bymaster et al., 2002). Treatments for schizophrenia also include antidepressants and benzodiazepines (Mueser & McGurk, 2004). However, many schizophrenic patients fail to take their medication due to the adverse side effects (Mueser & McGurk, 2004). In some cases, psychological treatment is also tried in addition to pharmacological treatments (Mueser & McGurk, 2004).

New treatments for schizophrenia are being employed using cholinergic agonists because the cholinergic system is known to modulate the dopaminergic system (Bymaster et al., 2002). Cholinergic agonists have been shown to have antipsychotic effects in schizophrenics (Andersen et al., 2003). In some cases, cholinesterase inhibitors have been administered in conjunction with the traditional antipsychotics with mixed results. Some schizophrenics see an improvement in cognitive symptoms whereas others see no apparent benefit (Raedler et al., 2007).

## **Binding and functional properties of xanomeline**

Xanomeline (3-(4-hexyloxy-1,2,5-thiadiazol-3-yl)-1,2,5,6-tetrahydro-1-methylpyridine) was developed by Eli Lilly as a potential therapeutic agent for the

treatment of AD (Sramek et al., 1995). Xanomeline is based on an arecoline derivative that is found naturally as the active ingredient in betel nuts (Raedler et al., 2007). In the initial analysis of xanomeline, it was found to have a high affinity for muscarinic receptors but lacked affinity at other non-muscarinic types of receptors (Bymaster et al., 1997; Mirza et al., 2003).

Xanomeline is a potent agonist that is able to bind to all subtypes of muscarinic receptors with similar potency, but is thought to be functionally selective for the M1 and M4 subtypes (Bymaster et al., 1997; Shannon et al., 1994). Functionally, xanomeline is a potent full agonist at the M1 receptor. In contrast, it is a potent partial agonist at the M2 and M5 receptors (Grant & El-Fakahany, 2005; Jakubik et al., 2006). Xanomeline has been shown to stimulate a functional response at the M1 receptor using multiple types of functional assays such as PI hydrolysis, citrulline formation and ERK phosphorylation in CHO cells (Christopoulos et al., 1999; Christopoulos et al., 1998) (Christopoulos, personal communication). At the M1 and M2 receptors, xanomeline exhibits selectivity in the coupling of the receptor to specific G proteins (Jakubik et al., 2006). Contradictory research on the functional properties of xanomeline conducted by Wood et al. (1999) determined that xanomeline was a full agonist at the M3 and M4 receptors, and a partial agonist at M1, M2 and M5 receptors. One possible explanation for the difference may be that there was receptor reserve at the M3 receptor, which made xanomeline appear to be a full agonist under these conditions (Wood et al., 1999). Additionally, they were using microphysiometry which detects changes in extracellular acidification as a measure of receptor activation (Wood et al., 1999).

Xanomeline has been shown to have a unique binding profile. Research conducted by Christopoulos et al (1998) at the M1 receptor demonstrated that there was not simple competition between xanomeline and atropine since the slope of the

functional PI assay curve was altered rather than simply shifted to the right as was observed with carbachol and atropine. Additionally, brief treatment with xanomeline resulted in a decrease in radioligand affinity in saturation binding experiments, also indicative of a non-competitive interaction (Christopoulos et al., 1998). Xanomeline was also shown to bind to the M1 receptor in a manner that was resistant to washout and was dependent on exposure time (Christopoulos et al., 1998). When xanomeline and atropine were present concurrently, atropine was able to block a functional response. However, when atropine was washed out xanomeline was again able to stimulate a functional response at the M1 receptor, suggesting some type of competitive interaction at the orthosteric site (Christopoulos et al., 1998). Furthermore, the presence of xanomeline in a competition binding assay with [<sup>3</sup>H]NMS, resulted in complete inhibition of [<sup>3</sup>H]NMS which is thought to be a result of competitive interactions with the receptor orthosteric site (Christopoulos et al., 1998). Additionally, there is also evidence that xanomeline interacts with the M1 receptor orthosteric site through the transmembrane domain of the receptor in a similar manner to that of carbachol, since the mutation of threonine 192 to alanine, results in a decrease in agonist affinity for both drugs compared to the affinity for wild-type receptors (Tejada et al., 2006). Thus it appears that xanomeline interacts with the receptor in both a competitive and non-competitive manner (Christopoulos et al., 1998).

### **Proposed mechanisms of xanomeline binding**

Based on the previous findings, a model for xanomeline binding at the M1 receptor was proposed. Binding of xanomeline at the M1 receptors occurs at both the orthosteric (primary) site (reversible binding) and an allosteric (secondary) site (wash-resistant binding) (Christopoulos et al., 1998). However, it is unclear if binding of

xanomeline occurs through the same mechanisms at other receptor subtypes. The model of xanomeline binding proposed by Christopoulos et al. (1998) was based on the interaction of the  $\beta_2$ -adrenoceptor agonist salmeterol studied by Coleman et al (1996) and Nials et al (1993).

### **Wash-resistant xanomeline binding**

Current laboratory research has primarily focused on the effects of xanomeline binding at the M1 receptor. Binding of xanomeline in a wash-resistant manner at the M1 receptor occurs rapidly (Jakubik et al., 2002). Wash-resistant binding also occurs at the M2 receptor, but the kinetics are slower than at the M1 (Jakubik et al., 2006). Additionally, research by Grant & El-Fakahany (2005) demonstrated that xanomeline is also able to bind in a wash-resistant manner at the M5 receptor. The binding of xanomeline to the M1 receptor occurs even if the primary binding site is blocked (Jakubik et al., 2002). The dissociation of the wash-resistant component of xanomeline binding from the M1 receptor occurs very slowly with a half life of over 30 hours (Jakubik et al., 2002). In addition, incubation of cells that express the M1 muscarinic receptor with xanomeline followed by washing leads to an apparent decrease in the number of binding sites as determined by saturation binding (Jakubik et al., 2002).

The alkyloxy side chain of xanomeline is necessary for the occurrence of wash-resistant binding to the M1 receptor (Jakubik et al., 2004). When the hydrophobic tail of xanomeline (hexyl, 6 carbons) is reduced to propyl (3 carbons) or methyl (1 carbon), the xanomeline analog no longer is able to bind in a wash-resistant manner at the M1 receptor (Jakubik et al., 2004; Kane et al., 2008). Furthermore, the xanomeline analogs with shortened side-chains that were still able to bind in a wash-resistant manner (butyl, 4 carbons and pentyl, 5 carbons) were able to interfere with wash-resistant binding of

xanomeline demonstrating that the analogs are binding to the receptor in a manner similar to that of xanomeline (Jakubik et al., 2004). Interestingly, although both the butyl and pentyl xanomeline analogs were able to bind in a wash-resistant manner only the pentyl analog was still able to stimulate a functional response in a wash-resistant manner (Jakubik et al., 2004).

Additional experiments were conducted to determine if xanomeline binds specifically in a wash-resistant manner to the receptor or if it simply partitions in a nonspecific manner into the lipid membrane. Xanomeline was able to bind in a wash-resistant manner when purified M1 receptors were reconstituted in liposomes, but not when receptors or liposomes were exposed to xanomeline alone prior to reconstitution. This demonstrates that xanomeline binds to the M1 receptor in a wash-resistant manner; however, the receptor needs to be in a lipid environment for this binding to take place (Jakubik et al., 2004). The importance of a lipid environment can also be seen by lowering the pretreatment temperature to 4 or 15°C which resulted in a decrease in affinity of wash-resistant xanomeline binding (Jakubik et al., 2004).

Jakubik et al (2004) suggested that if the underlying mechanism of wash-resistant binding of xanomeline is elucidated it may lead to the development of both muscarinic agonists and antagonists with long duration of action and perhaps better pharmacological selectivity. In order to determine the mechanism of xanomeline binding it is important to investigate the action of xanomeline across subtypes using the same receptor expression system.

### **Xanomeline as a treatment for Alzheimer's disease and schizophrenia**

Although the mechanisms for Alzheimer's disease and schizophrenia are likely different, there is an overlap in some of the symptoms, such as delusions, hallucinations,

memory deficits, etc., which can be alleviated by treatment with cholinergic agonists (Mirza et al., 2003). Thus, research has been conducted using xanomeline as a potential therapeutic for both diseases in animal models and in clinical studies.

### **Xanomeline treatment in Alzheimer's disease models**

The characteristics of xanomeline binding and function were studied in animal models. Xanomeline was found to bind with high affinity in rat hippocampus and rabbit vas deferens, both of which are thought to primarily express M1 receptors (Shannon et al., 1994). In contrast, minimal binding of xanomeline was observed in guinea pig atria (M2 receptors) or guinea pig urinary bladder (M3 receptors) (Shannon et al., 1994). Treatment with xanomeline resulted in an increase in striatal levels of a metabolite of dopamine and an increase in the levels of acetylcholine in the rat, suggesting that both dopamine and acetylcholine release were elevated in response to xanomeline. Since there is known to be an interaction between dopamine and muscarinic receptors and xanomeline has not been shown to interact with dopamine receptors, it is thought that the response is mediated by M1 receptors (Bymaster et al., 1994). Interestingly, the effects of xanomeline on these neurotransmitters in other brain regions were variable (Bymaster et al., 1994). When rats were treated with xanomeline, an increase in salivation, which is thought to be mediated by the M3 receptor, was only detected in one of five animals at a high dose (Bymaster et al., 1994). When tissue slices of rat brain were treated with xanomeline *in vitro* the release of acetylcholine was inhibited following additional stimulation, but the basal release was unchanged (Machova et al., 2007). Furthermore, when tissues were preincubated with xanomeline there was a concentration-dependent decrease in acetylcholine release following carbachol-induced stimulation (Machova et al., 2007). These results are most likely due to the activation of

the M2 and/or M4 muscarinic receptors. Taken together these results suggest that the actions of xanomeline may be efficacious in the treatment of AD.

### **Clinical trials with xanomeline for the treatment of Alzheimer's disease**

Clinical trials were conducted to determine the effects of xanomeline treatment in AD. Initial data demonstrated that xanomeline was able to cross the blood brain barrier, and thus would be able to reach the targeted M1 receptors (Bymaster et al., 1997). Interestingly, the concentration of xanomeline in the brain was found to be ten to twenty times higher compared to plasma levels (Mirza et al., 2003). Xanomeline was also shown to undergo high first pass metabolism, which occurs through  $\beta$ -oxidation of the aliphatic side chain and N-oxidation of the tetrahydropyridine ring (Bymaster et al., 1997). The N-desmethyl metabolites have been shown to be pharmacologically active (Mirza et al., 2003).

Xanomeline was able to reduce cognitive deficits and behavioral symptoms in Alzheimer's disease patients, which are thought to be due to activation of the M1 receptor (Bymaster et al., 1997). This occurred in patients that had both mild and moderate AD (Bymaster et al., 1997). The maximum effect on cognition occurred between eight and ten weeks after the start of drug administration (Bodick et al., 1997 (b); Bodick et al., 1997 (a)). In addition to improvements in cognitive deficits, treatment of AD patients with xanomeline resulted in a reduction in outbursts, suspiciousness, delusions, agitation and hallucinations (Bodick et al., 1997 (b); Mirza et al., 2003). However, treatment with xanomeline also resulted in adverse side effects such as hypersalivation, bradycardia, gastrointestinal tract problems, sweating, nausea, vomiting, and syncope, which are thought to be mediated mainly by non-selective activation of the M2 and M3 muscarinic receptors (Bodick et al., 1997 (b); Bymaster et al., 1997). In a



clinical trial conducted by Sramek et al. (1995) 94 percent of patients experienced some side effects. Interestingly, higher plasma concentrations of xanomeline appeared to correlate with more adverse effects (Sramek et al., 1995). The side effects of xanomeline may be due to non-selective receptor activation by xanomeline and/or as a result of activation by metabolites from the metabolism of the alkyloxy side chain of xanomeline (Messer, 2002). Additionally, it may be possible that xanomeline has different effects in different areas in the brain (Mirza et al., 2003).

### **Xanomeline treatment in animal models of schizophrenia**

Startle amplitude, locomotor activity, prepulse inhibition and conditioned avoidance are used as behavioral methods to assess schizophrenia in animal models (Bymaster et al., 2002). The administration of xanomeline resulted in a decrease in startle amplitude in a dose-dependent manner (Jones et al., 2005; Stanhope et al., 2001). Treatment with a non-selective muscarinic agonist, pilocarpine, was shown to have a similar effect compared to xanomeline on decreasing startle amplitude, demonstrating that the decrease in response is mediated by muscarinic receptors (Stanhope et al., 2001). In animals, treatment with amphetamine has been shown to increase locomotion as a result of an increase in dopamine in the synapse (Bymaster et al., 2002). Administration of xanomeline was shown to decrease amphetamine-induced hyperlocomotion, while having no effect on baseline locomotion (Bymaster et al., 2002; Stanhope et al., 2001). Schizophrenics have been shown to have a decrease in prepulse inhibition of the acoustic startle reflex (Raedler et al., 2007). In prepulse inhibition, a nonstartling tone that occurs just prior to an intense stimulus reduces the startle response (Bymaster et al., 2002). Treatment with xanomeline is able to reverse this deficit in an animal model of schizophrenia (Raedler et al., 2007; Stanhope et al.,

2001). Taken together these animal models suggest that xanomeline may alleviate some of the symptoms of schizophrenia in humans. The antipsychotic profile of xanomeline may be mediated through the M1 or M4 muscarinic receptor, which at this time remains unclear (Mirza et al., 2003).

### **Clinical trials with xanomeline for the treatment of schizophrenia**

A recent clinical trial was conducted employing xanomeline as a treatment for schizophrenia. Treatment of schizophrenic patients with xanomeline resulted in improvements in total Positive and Negative Syndrome Scale scores, verbal learning and short-term memory function by the end of the first week of a four week treatment period (Shekhar et al., 2008). There were minimal adverse side effects such as gastrointestinal tract problems, but they were not severe enough to discontinue treatment (Shekhar et al., 2008). Thus, preliminary results suggest that xanomeline may be beneficial in the treatment of schizophrenia.

### **Summary**

Muscarinic receptors (M1-M5) are G protein coupled receptors that are located widely throughout the brain and periphery. These receptors are known to be involved in a variety of diseases, such as Alzheimer's disease and schizophrenia, which has made them targets for drug development. Currently, subtype selective agonists and antagonists that bind at the orthosteric site are not available due to the highly conserved nature of the orthosteric site. New research is beginning to focus on allosteric sites as drug targets because these sites are thought to be less conserved across subtypes, thus potentially resulting in greater subtype selectivity. Xanomeline is a novel muscarinic receptor agonist that binds at both the orthosteric site and at an allosteric site. This

thesis will address the effects of acute and long-term xanomeline treatment on muscarinic receptors in terms of binding and functional properties.

## **CHAPTER 2: Materials and methods**

## **MATERIALS AND GENERAL TREATMENT PROTOCOLS**

### **Materials**

[<sup>3</sup>H]N-Methylscopolamine ([<sup>3</sup>H]NMS) (81 Ci/mmol) was purchased from DuPont (Wilmington, DE); [<sup>3</sup>H]quinuclidinyl benzilate ([<sup>3</sup>H]QNB) (52 Ci/mmol) was purchased from DuPont (Wilmington, DE); *myo*-[<sup>3</sup>H]inositol (71 Ci/mmol) was purchased from GE Healthcare (Little Chalfont, Buckinghamshire, UK); [<sup>14</sup>C]inositol-1-phosphate (300 mCi/mmol) was purchased from American Radiolabeled Chemicals (St. Louis, MO); Dulbecco's modified Eagle's medium (DMEM) was purchased from Invitrogen (Carlsbad, CA); geneticin was obtained from Calbiochem (San Diego, CA); siRNA was purchased from Ambion (Austin, TX); and bovine calf serum was supplied by Hyclone Laboratories (Logan, UT). Xanomeline tartrate was a generous gift from Eli Lilly & Co. (Indianapolis, IN); all other reagents were purchased from Sigma-Aldrich (St. Louis, MO).

### **Cell culture**

Chinese hamster ovary (CHO) cells stably transfected with the human M1-M4 muscarinic acetylcholine receptors (hM1-hM4) were provided by Dr. M. Brann, University of Vermont Medical School. The genes encoding the rat M1 wild-type (rM1) and mutant (rM1<sup>123</sup>) muscarinic receptors were stably expressed in CHO cells (Zhu et al., 1991). All cells were grown in Dulbecco's modified Eagle's medium supplemented with 10% bovine calf serum and 50 µg/ml geneticin. Cells were grown to confluency in 24-well plates, culture flasks or 10-cm dishes at 37°C in a humidified atmosphere consisting of 5% CO<sub>2</sub> / 95% air.

### **Pretreatment with agonist**

CHO cells expressing hM1-hM4 receptors were incubated with agonists in culture media at 37°C for one minute or one hour, then washed three times with HEPES buffer (110 mM NaCl, 5.4 mM KCl, 1.8 mM CaCl<sub>2</sub>, 1 mM MgSO<sub>4</sub>, 25 mM glucose, 20 mM HEPES, 58 mM sucrose; pH 7.4 ± 0.02; 340 ± 5 mOsm) to remove any unbound drug. Cells were used immediately or allowed to incubate for 23-24 hours in ligand-free media. Another group of cells was incubated with agonist in culture media for 24 hours before washing three times with HEPES buffer. All pretreatments and radioligand binding assays were performed on cells in monolayer at 37°C unless otherwise noted.

### **siRNA treatment**

CHO cells expressing hM3 receptors were incubated with 50 nM siRNA each for G<sub>q</sub> and G<sub>11</sub> G-proteins in Opti-Mem for 6-7 hours at 37°C. Culture medium with no geneticin was added and the cells were allowed to incubate overnight. Subsequent pretreatment with xanomeline was the same as stated under pretreatment with agonist.

### **Determination of protein content**

Protein determinations were performed according to the method of Bradford (Bradford, 1976) following agonist pretreatment. The changes in protein content following agonist treatments were used to correct radioligand binding curves as necessary. Briefly, cells were incubated with carbachol (100 µM – 10 mM), pilocarpine (1 µM – 100 µM), xanomeline (1 µM – 100 µM) or DMSO (0.01 – 1 %) for one hour in HEPES buffer followed by washing, which corresponds to the continuous presence of the agonist in the binding assay. Additional cells were treated with carbachol (100 µM – 10 mM), pilocarpine (1 µM – 100 µM), xanomeline (1 µM – 100 µM) or DMSO (0.01 – 1

%) for one minute or one hour in media followed by washing and used immediately or subsequent to 23-24 hour incubation in control media. Another group of cells was treated with agonists or DMSO for 24 hours prior to washing. In all cases reactions were stopped by incubation with 1 M NaOH for 30 minutes. Samples were either used directly or following dilution in the Bradford assay.

## **RADIOLIGAND BINDING ASSAYS**

### **Time course**

Experiments were conducted to determine the time course of xanomeline wash-resistant binding. CHO hM1-hM4 cells were incubated for various times (< 15 seconds to 30 minutes) with a single concentration of xanomeline (10  $\mu$ M) and then washed three times with HEPES buffer. In the binding assay, cells were incubated with the muscarinic receptor ligand [ $^3$ H]N-methylscopolamine ([ $^3$ H]NMS) (0.2 nM) for one hour at 37°C. In all instances, non-specific binding was determined using 10  $\mu$ M atropine during the binding assay. Incubations were terminated by removing the buffer containing free radioligand and then cells with bound radioligand were dissolved in 1M NaOH. Bound radioactivity (disintegrations per minute (dpm)) was quantitated by liquid scintillation spectrometry.

### **Inhibition of [ $^3$ H]NMS binding following agonist pretreatments**

This protocol was employed to assess agonist potency. In all cases, the IC<sub>50</sub> (concentration of xanomeline that resulted in the inhibition of 50 percent of radioligand binding) was used as a measure of the potency of xanomeline binding. CHO hM1-hM4 cells were incubated concurrently with 0.2 nM [ $^3$ H]NMS and increasing concentrations of xanomeline (1 nM – 100  $\mu$ M), carbachol (10 nM – 10 mM) or pilocarpine (1 nM – 100  $\mu$ M) for one hour. In order to assess the effects of prolonged agonist treatments, cells

were pretreated with increasing concentrations of xanomeline (1 pM – 100 μM), carbachol (1 pM – 10 mM) or pilocarpine (1 pM – 100 μM) for one minute or one hour followed by washing three times with HEPES buffer and used immediately or after incubation for 24 or 23 hours, respectively, in ligand free media. An additional group of cells was pretreated with agonists for 24 hours prior to washing three times with HEPES buffer. Subsequently, cells were incubated with [<sup>3</sup>H]NMS (0.2 nM) for one hour. In all instances, pretreatment and radioligand binding were performed in monolayer at 37°C. Non-specific binding was determined using 10 μM atropine. Incubations were terminated as described above. Protein determinations were performed according to the method of Bradford (Bradford, 1976) as described above. Bound radioactivity (dpm) was quantitated by liquid scintillation spectrometry.

Due to the fact that xanomeline was dissolved in dimethyl sulfoxide (DMSO), additional control experiments were performed to ensure that DMSO did not contribute to the observed effects of xanomeline. Radioligand binding and protein content assays were conducted as described above using DMSO concentrations corresponding to those in contact with cells during the various treatment modalities with xanomeline (0.01 – 1%).

Further experiments were conducted at the hM1 and hM3 receptors that were the same as stated above except saturating concentrations of the muscarinic receptor ligands [<sup>3</sup>H]NMS (2.9 nM) or [<sup>3</sup>H]QNB (1.4 nM) were used. The concentrations of the [<sup>3</sup>H]NMS and [<sup>3</sup>H]QNB were chosen to achieve 93 percent fractional receptor occupancy.



### **[<sup>3</sup>H]NMS saturation binding**

A saturation binding paradigm was used to determine the total cell-surface receptor density ( $B_{max}$ ) and radioligand-receptor equilibrium dissociation constant ( $K_d$ ). CHO hM1-hM4 cells were incubated in monolayer for one hour at 37°C in the absence or presence of xanomeline (10  $\mu$ M), carbachol (10  $\mu$ M; hM3 receptor only) or pilocarpine (10  $\mu$ M; hM3 receptor only) followed by three washes with HEPES buffer and used immediately or after 23 hour incubation in control media. An additional group of cells was treated with xanomeline or carbachol (hM3 receptor only) for 24 hours before washing three times with HEPES buffer. Subsequently, cells were incubated in monolayer with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 8.9 nM) for one hour at 37°C. Non-specific binding was determined using 10  $\mu$ M atropine. Reactions were terminated and bound radioactivity was quantitated as described above. Protein determinations were performed according to the method of Bradford (Bradford, 1976) described above.

### **FUNCTIONAL ASSAYS**

#### **Assay of phosphoinositide (PI) hydrolysis**

CHO hM3 or hM1 cells in 24-well plates were loaded with *myo*-[<sup>3</sup>H]inositol (1  $\mu$ Ci/ml) at 37°C 24 hours prior to being subjected to several different treatment protocols with agonists and determination of inositol phosphate production.

1) Continuous presence of agonist: Labeled untreated cells were incubated in the presence of increasing concentrations of xanomeline (10 pM – 10  $\mu$ M), pilocarpine (100 nM – 60  $\mu$ M), or carbachol (10 nM – 100  $\mu$ M) in the presence of 10 mM LiCl for one hour at 37°C.

2) Acute agonist treatment: Cells were pretreated with increasing concentrations of xanomeline (100 pM – 100  $\mu$ M) for one minute or one hour, at which time cells were washed three times with HEPES buffer. In other cases, cells were treated with a single high concentration of carbachol (10  $\mu$ M) or pilocarpine (10  $\mu$ M) for one hour followed by washing. Cells were then incubated for one hour at 37°C in the presence of 10 mM LiCl but in the absence of further agonist stimulation.

3) Xanomeline treatment followed by stimulation with a second agonist: Cells were pretreated for one minute or one hour with a single concentration of xanomeline (10  $\mu$ M for hM3; 300 nM for hM1), washed three times with HEPES buffer then used immediately or incubated in control media for 24 or 23 hours, respectively. Additional cells were pretreated with xanomeline (10  $\mu$ M hM3; 300 nM hM1) for 24 hours, washed and used immediately in the functional assay. Subsequently, cells were incubated with increasing concentrations of carbachol (10 nM – 10 mM), pilocarpine (10 nM – 1 mM; hM3 only) or xanomeline (1 nM - 100  $\mu$ M; hM3 only) for one hour in the presence of 10 mM LiCl.

4) Agonist treatment followed by stimulation with a second agonist at the hM3 receptor: Cells were pretreated for one hour with a single concentration of carbachol (10  $\mu$ M) or pilocarpine (10  $\mu$ M), washed three times with HEPES buffer and used immediately or after 23 hour incubation in control media. Additional cells were pretreated with carbachol (10  $\mu$ M) or pilocarpine (10  $\mu$ M) for 24 hours prior to washing three times with HEPES buffer. Subsequently, cells were incubated for one hour with carbachol (10 nM – 10 mM), pilocarpine (10 nM – 1 mM) or xanomeline (1 nM - 100  $\mu$ M) in the presence of 10 mM LiCl.

5) Time course of agonist stimulation: The time course of xanomeline-induced persistent activation and antagonism of carbachol-elicited PI hydrolysis was established.

Cells were exposed to xanomeline (10  $\mu$ M for hM3; 300 nM for hM1) for increasing amounts of time (30 minutes – 24 hours) followed by washing and immediate use in the functional assay. Alternatively, cells were treated with xanomeline (10  $\mu$ M for hM3; 300 nM for hM1) for one hour followed by extensive washing and incubation in control media for increasing lengths of time (1 – 23 hours) as indicated in the Results. In both cases, cells were incubated for one hour in a LiCl-containing buffer in the absence of further agonist stimulation or in the presence of a single high concentration of carbachol (10  $\mu$ M).

In all cases, reactions were stopped with 0.3 M HClO<sub>4</sub> and neutralized with 0.15 M K<sub>2</sub>CO<sub>3</sub>. The samples were centrifuged (1500 x g; 15 min) and total inositol phosphates in the supernatant were separated by ion exchange chromatography (AG1-X8 resin). In all cases, [<sup>14</sup>C]inositol-1-phosphate was used as an internal recovery standard. Radioactivity (dpm) was quantitated by liquid scintillation spectrometry and adjusted for <sup>14</sup>C recovery.

Control experiments were conducted using CHO hM1 and hM3 cells to insure that the pool of [<sup>3</sup>H]*myo*-inositol was not depleted following xanomeline pretreatments. To this accord, cells were pretreated with xanomeline as detailed above and then incubated with 2.5 U of exogenous phospholipase C in the presence of 10 mM LiCl. The reactions were stopped and radioactivity quantitated as described above.

Alternatively, CHO cells expressing the hM1 receptor were treated in monolayer but the functional assay was performed in suspension to determine the effect of xanomeline pretreatment on the ability of agonists to further stimulate the production of inositol phosphates. CHO hM1 cells were incubated in monolayer with *myo*-[<sup>3</sup>H]inositol

(1  $\mu\text{Ci/ml}$ ) for 24 hours at 37°C prior to being subjected to the following different treatment protocols with agonists and determination of inositol phosphate production.

1) Xanomeline treatment followed by agonist stimulation: Cells were pretreated for one minute or one hour with a single concentration of xanomeline (300 nM), washed three times with HEPES buffer and then used immediately or incubated in control media for 24 or 23 hours, respectively. Additional cells were pretreated with xanomeline (300 nM) for 24 hours prior to washing. Cells were then harvested by trypsinization, centrifuged, and washed three times in HEPES buffer. Subsequently, cells were distributed to assay tubes (500,000 cells /assay tube), and allowed to incubate for 10 minutes at 37°C. Concentration-response curves for the stimulation of inositol phosphate production were determined for carbachol (100 nM – 10 mM), pilocarpine (10 nM – 1 mM) or xanomeline (1 pM – 10  $\mu\text{M}$ ) in the presence of 10 mM LiCl.

2) Carbachol treatment followed by agonist stimulation: Cells were pretreated for 24 hours with a single concentration of carbachol (10  $\mu\text{M}$ ) prior to washing. Cells were then harvested as described above. Subsequently, cells were incubated for one hour with carbachol (10 nM – 1 mM) or xanomeline (10 pM – 100  $\mu\text{M}$ ) in the presence of 10 mM LiCl.

In all cases, the reaction was stopped with chloroform/methanol (2:1) and centrifuged at 450 x g for 15 min. For all experiments, [ $^{14}\text{C}$ ]inositol-1-phosphate was added to each sample as an internal recovery standard. Total inositol phosphates were separated by ion exchange chromatography (AG1-X8 resin). Radioactivity (dpm) in each sample was quantitated by liquid scintillation spectrometry and adjusted for  $^{14}\text{C}$  recovery.

## **VARIATIONS ON RADIOLIGAND BINDING AND FUNCTIONAL ASSAY PARADIGMS**

### **Atropine blockade of receptor activation using saturation binding and functional assays**

CHO hM3 cells were grown in 24-well plates and pretreated in monolayer for one hour at 37°C in the absence or in the presence of xanomeline (10 µM) alone or in conjunction with atropine (10 µM) followed by three washes with HEPES buffer and 23 hour incubation in control media. A second group of cells was untreated or treated with xanomeline for one hour, washed extensively and then treated with atropine for 23 hours. In the binding assay, cells were incubated in monolayer with increasing concentrations of [<sup>3</sup>H]NMS (0.01 - 13.5 nM) for one hour at 37°C. Non-specific binding was determined using 10 µM atropine. Reactions were terminated and radioactivity quantitated as described above. In the functional assay cells were incubated with increasing concentrations of carbachol (10 nM – 10 mM) for one hour in the presence of 10 mM LiCl. The reaction was stopped and radioactivity (dpm) quantitated as described above.

CHO hM1 cells were grown in tissue culture flasks. Pretreatments were performed in monolayer at 37°C in the absence or in the presence of xanomeline (3 µM saturation; 300 nM function) alone or in conjunction with atropine (10 µM) followed by three washes with HEPES buffer and 23 hour incubation in control media. A second group of cells was untreated or treated with xanomeline for one hour, washed extensively and then treated with atropine for 23 hours. Cells were then harvested by trypsinization, followed by centrifugation (300 x g, 3 min) and re-suspension of the pellet in HEPES buffer (three times). In the suspension binding assay, cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.02 to 6 nM) for one hour at 37°C using 100,000 cells/assay tube in a total volume of 1 ml. Nonspecific binding was determined

using 10  $\mu\text{M}$  atropine. The reaction was terminated by filtration on Whatman GF/C filters (Whatman Schleicher and Schuell, Keene, NH) using a Brandel cell harvester (Brandel Inc., Gaithersburg, MD). Filters were washed three times with 4-ml aliquots of ice-cold saline and dried before radioactivity (dpm) was quantitated using liquid scintillation spectrometry. In the functional assay, 500,000 cells/assay tube were allowed to incubate for 10 minutes at 37°C prior to the addition of carbachol (100 nM – 1 mM). The reaction was allowed to proceed for one hour in the presence of 10 mM LiCl before being stopped with chloroform/methanol (2:1). The remainder of the procedure was the same as described above for the PI assay in suspension.

### **Block of receptor activation using a receptor mutant in binding and functional assays**

Experiments were conducted to determine the role of receptor function on the long-term effects of xanomeline at the rM1 receptor. CHO rM1 or rM1<sup>123</sup> mutant cells were exposed to increasing concentrations of carbachol (10 nM – 500  $\mu\text{M}$ ) or xanomeline (10 pM – 100  $\mu\text{M}$ ) in suspension to determine the functional response. The PI assay was performed as noted previously for cells in suspension. Subsequently, inhibition of [<sup>3</sup>H]NMS and saturation binding assays were performed. Briefly, in the inhibition of [<sup>3</sup>H]NMS binding assay, cells were treated concurrently with 0.2 nM [<sup>3</sup>H]NMS and increasing concentrations of xanomeline (100 pM – 100  $\mu\text{M}$ ) for one hour. Additional cells were treated with xanomeline (1 pM – 100  $\mu\text{M}$ ) for one hour followed by washing and used immediately or following 23 hour incubation in control media. An additional group of cells was treated with xanomeline for 24 hours prior to washing. The inhibition of [<sup>3</sup>H]NMS binding assay and radioligand quantification was performed as described previously for binding in suspension. In the [<sup>3</sup>H]NMS saturation binding

experiments rM1 or rM1<sup>123</sup> mutant cells were exposed to xanomeline (3  $\mu$ M) for one hour followed by three washes with HEPES buffer and used immediately or subsequent to 23 hour incubation in the absence of free ligand. An additional group of cells was treated with xanomeline for 24 hours prior to washing. The radioligand binding assay and quantification were performed as previously described for suspension binding. Control experiments utilizing DMSO in [<sup>3</sup>H]NMS binding and protein assays were conducted as described above to determine the effects of agonist pretreatments on rM1 or rM1<sup>123</sup> mutant receptors.

### **Use of siRNA blockade of receptor function in saturation binding and functional assays**

Experiments were conducted to determine the role of receptor function on the long-term effects of xanomeline at the M3 receptor. siRNA transfections were performed as noted above. The transfection protocol was unsuccessful in cells seeded in 24-well plates. Therefore, transfections were performed in 10-cm dishes. To confirm a decrease in inositol phosphates production, non-transfected and transfected cells were removed from the 10-cm dishes by trypsinization and centrifuged (300 x g, 2 min) and re-suspended three times in HEPES buffer. Subsequently, in the functional assay in suspension, cells were incubated with carbachol (1  $\mu$ M and 100  $\mu$ M) or xanomeline (100 nM and 10  $\mu$ M) for one hour at 37°C. The reaction was stopped and the accumulation of inositol phosphates was determined as previously stated for suspension.

Western blot analysis was performed to confirm a decrease in G<sub>q/11</sub> G protein expression following siRNA pretreatment. Briefly, cells were washed and centrifuged. The pellet was reconstituted in 250  $\mu$ l of RIPA lysis buffer and placed on ice for 20 minutes. Cells were passed through a 23-gauge needle to further homogenize the

sample, then incubated on ice for 20 minutes and spun at 14,000 x g for 20 minutes at 4°C. The supernatant was collected and the amount of protein in the sample was determined using the method of Bradford (Bradford, 1976). Fifty µg of protein from each sample was loaded onto a 12.5% SDS polyacrylamide gel. The proteins were separated by electrophoresis then transferred onto a nitrocellulose membrane, which was blocked overnight at 4°C with a dry milk solution. The G<sub>q/11</sub> G proteins were detected with a rabbit anti-G<sub>q/11</sub> antibody. Horseradish peroxidase-labeled anti-rabbit secondary antibody was used. The membrane was then treated with chemiluminescence reagents and exposed on x-ray film. The approximately 45 kD band was quantified using a Bio-Rad densitometer. The optical density of each sample was measured followed by subtraction of background.

Subsequently, in the [<sup>3</sup>H]NMS saturation binding assay, non-transfected or siRNA transfected cells were incubated for one hour at 37°C in the absence or presence of xanomeline (10 µM) followed by washing and immediate use or further incubation for 23 hours in control medium. Additional cells were treated with xanomeline for 24 hours prior to washing three times with HEPES buffer. Cells were then removed from the 10 cm dishes by trypsinization, centrifuged (300 x g, 2 min) and re-suspended in HEPES buffer (three times). Subsequently, cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.04 – 8.5 nM) in suspension for one hour at 37°C. The remaining details are the same as previously described for saturation binding in suspension.

### **[<sup>3</sup>H]NMS saturation binding following propyl benzyl choline mustard (PrBCM) treatment**

Experiments were conducted to determine the effect of decreasing the level of cell-surface receptor density on saturation binding parameters using an irreversible



receptor alkylating agent. The levels of cell-surface receptor density at the M4 receptor were decreased to the levels observed at the M2 receptor to determine if a lower level of receptor density had an influence on the long-term effects of xanomeline binding. To determine the concentration of propylbenzyl choline mustard (PrBCM) that resulted in an approximately 50 percent decrease in receptor expression CHO cells expressing the hM4 receptor were treated with increasing concentrations of 0.1 – 10 nM cyclized PrBCM (Hu & El-Fakahany, 1990) for 15 minutes, followed by washing three times with HEPES buffer. Subsequently, cells were exposed to 6.5 nM [<sup>3</sup>H]NMS for one hour at 37°C. The reaction was stopped and radioligand binding was determined as described above for binding assays in monolayer. Subsequent saturation binding experiments were conducted using 2 nM PrBCM. In this case cells were exposed to 2 nM PrBCM for 15 minutes followed by three washes with HEPES buffer. Cells were then pretreated with xanomeline (10 μM) for one hour followed by washing and waiting 23 hours or treated with xanomeline for 24 hours prior to washing. Subsequently, untreated or xanomeline pretreated cells were exposed to increasing concentrations of [<sup>3</sup>H]NMS (0.03 – 7 nM) for one hour at 37°C. The rest of the protocol for the termination of the reaction and radioligand quantitation is as described above for saturation binding assays in monolayer.

### **[<sup>3</sup>H]NMS saturation binding following pretreatment with pertussis toxin**

Experiments were conducted to determine the effect of blocking the G<sub>i/o</sub> G proteins on saturation binding parameters at the hM4 receptor. Cells were treated with pertussis toxin (100 ng/ml) in media for 24 hours. Subsequently, cells were pretreated with xanomeline (10 μM) for one hour followed by washing three times with HEPES buffer and further incubation for 23 hours in ligand free media containing pertussis toxin.

Additional cells were treated concurrently with xanomeline and pertussis toxin for 24 hours prior to washing. Cells were then treated with increasing concentrations of [<sup>3</sup>H]NMS (0.02 – 7.5 nM) for one hour at 37°C. The reaction was stopped and radioligand binding determined as described above for saturation binding on cells in monolayer.

### **DATA ANALYSIS**

All analyses were conducted using Prism 4.0 (GraphPad Software Inc., San Diego, CA). Inhibition of [<sup>3</sup>H]NMS binding isotherms were analyzed via nonlinear regression to derive estimates of IC<sub>50</sub> (midpoint location or potency parameter). Data were refitted according to both one- and two-site mass-action binding models, and the better model was determined by an extra sum-of-squares test. The quasi-irreversible nature of xanomeline binding did not permit transforming IC<sub>50</sub> values to inhibition constants because such transformation assumes reversible competitive interaction. Saturation binding curves were analyzed via nonlinear regression to derive individual estimates of  $B_{\max}$  (total cell-surface receptor density) and  $K_d$  (radioligand-receptor equilibrium dissociation constant). PI hydrolysis isotherms were analyzed via non-linear regression. Data shown are the means ± standard error of the mean. Comparisons between mean values were performed by paired *t* tests, unpaired *t* tests or one way analysis of variance (ANOVA), as appropriate. A probability (*p*) value < 0.05 was taken to indicate statistical significance.

**CHAPTER 3: Determination of the acute and long-term effects of xanomeline binding at M1-M4 muscarinic receptors**

## **Introduction**

Muscarinic acetylcholine receptors (mAChR) are part of the G protein-coupled receptor superfamily. There are five subtypes of muscarinic receptors (M1-M5) (Caulfield & Birdsall, 1998). These receptors are expressed widely throughout the brain and periphery (Caulfield & Birdsall, 1998). Each receptor has a different primary physiological function with the M1 receptor involved in learning and memory (Eglen, 2005). M2 and M4 receptors are mainly located presynaptically and mediate autoinhibition of acetylcholine release in the hippocampus and cerebral cortex, respectively (Wess et al., 2003). M3 receptors are involved in salivation, food intake, and gastrointestinal tract motility and secretion (Eglen, 2005). M5 receptors facilitate dopamine release in the striatum (Wess et al., 2003). There is a high degree of homology between the orthosteric (primary) binding region across subtypes (Eglen, 2005), which makes it difficult to develop an agonist that binds selectively to a particular subtype (Wess, 2004). However, recent research suggests that there are secondary binding domains (allosteric binding sites) on muscarinic receptors that are less conserved across subtypes, which may lead to the development of agonists that are able to act in a more selective manner (Eglen, 2005). Prolonged exposure of receptors to an agonist can lead to changes in the binding properties of the receptor. These changes in binding properties can result from receptor internalization, down-regulation or uncoupling from the G protein (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990).

Xanomeline is a potent agonist that is able to bind to all subtypes of muscarinic receptors with similar potency (Shannon et al., 1994). However, it is thought to have functional selectivity for the M1 and M4 receptors (Bymaster et al., 1997; Shannon et al., 1994). However, it should be noted that the previous research was conducted using different receptor expression systems. Additionally, xanomeline binding at the M1

receptor has been shown to take place both at the orthosteric (primary) binding site in a reversible manner and at an allosteric (secondary) site in a wash-resistant manner (Christopoulos et al., 1998). However, it is unclear if binding of xanomeline occurs through the same mechanisms at other receptor subtypes.

We were interested in determining the short- and long-term effects of xanomeline binding across receptor subtypes. It is important to compare the effects of xanomeline binding in the same expression systems to avoid complications caused by factors not related to the type of muscarinic receptor expressed. Thus, experiments were conducted in Chinese hamster ovary (CHO) cells expressing M1-M4 muscarinic receptors to test the hypotheses that 1) xanomeline binds to all receptor subtypes in both a reversible and wash-resistant manner, 2) xanomeline wash-resistant binding results in long-term effects similar to those seen during prolonged treatment with agonists, such as internalization and down-regulation and 3) there is subtype selectivity in the long-term effects of wash-resistant xanomeline binding.

## **Results**

### **Time course of development of wash-resistant xanomeline binding at M1-M4 receptor subtypes**

The time course of xanomeline binding in a wash-resistant manner was determined at each of the muscarinic receptor subtypes expressed in CHO cells by measuring the decrease in [<sup>3</sup>H]NMS binding over time following xanomeline exposure and extensive washing. [<sup>3</sup>H]NMS binding in the absence of xanomeline was used as a control. In all cases, there was an immediate decrease in [<sup>3</sup>H]NMS binding when cells were briefly treated (< 15 seconds) with 10 μM xanomeline (Fig. 1). However, the decrease in [<sup>3</sup>H]NMS binding following xanomeline treatment for <15 seconds was

significantly ( $p < 0.05$ ) greater for the M1 (48%) and M3 (41%) receptor subtypes, than the M2 (24%) or M4 (30%) subtypes, suggesting that xanomeline wash-resistant binding occurs more rapidly at the former subtypes. Subsequently, there was a slower phase of inhibition of [ $^3\text{H}$ ]NMS binding that reached a plateau by approximately 10 minutes for all subtypes. Once again, the maximal decrease in [ $^3\text{H}$ ]NMS binding was significantly ( $p < 0.05$ ) greater at the M1 (81%) and M3 (83%) receptors, compared to the M2 (66%) and M4 (71%) receptors (Fig. 1). The results at the M1 receptor are in agreement with what Jakubik et al. (2002, 2006) observed. However, there is a discrepancy regarding the time course of xanomeline binding at the M2 receptor. Specifically, Jakubik et al. (2006) observed no decrease in [ $^3\text{H}$ ]NMS binding following treatment with xanomeline for <15 seconds and a more rapid decrease in the subsequent phase. These differences could be the result of variations in the experimental protocols employed. In the current protocol, the binding assay was performed on intact cells in monolayer whereas the protocol used by Jakubik et al. (2006) for the binding assay was one in which treated cells had been homogenized to create membrane preparations for the binding assay. Additionally, they used a higher concentration of [ $^3\text{H}$ ]NMS (1 nM), which could influence the ability of xanomeline to compete with and inhibit radioligand binding.

### **Differential effects of acute xanomeline treatment on inhibition of [ $^3\text{H}$ ]NMS binding at M1-M4 receptor subtypes**

Previous research from our laboratory and others has demonstrated that xanomeline binding occurs in a wash-resistant manner at the M1 receptor (Christopoulos et al., 1998; Grant & El-Fakahany, 2005; Jakubik et al., 2002). The current set of experiments were undertaken to compare the short-term effects of xanomeline binding across the M1-M4 receptor subtypes. The presence of xanomeline in the binding assay

medium caused a concentration-dependent and complete inhibition of [<sup>3</sup>H]NMS binding at all receptor subtypes (Fig. 2). The rank order of xanomeline potencies was M3 ≈ M4 > M2 > M1. When cells were treated with xanomeline for one minute or one hour followed by washing and immediate use there was a significant ( $p < 0.05$ ) decrease in xanomeline potency for all subtypes compared to the experimental protocol described above (Fig. 2; Table 1). Pretreatments for one minute, but not for one hour, resulted in incomplete inhibition of [<sup>3</sup>H]NMS binding at all four receptor subtypes (Table 1). For the brief one minute xanomeline pretreatments, the rank order of potencies of xanomeline was M1 > M2 > M3 ≈ M4. However, there was no difference in the potencies when the treatment time with xanomeline was increased to one hour.

Since xanomeline was dissolved in dimethylsulfoxide (DMSO), control experiments were conducted to ascertain if the vehicle contributed to any of the observed inhibition of [<sup>3</sup>H]NMS binding. The concentrations of DMSO used corresponded to those employed in the preparation of the various xanomeline concentrations. DMSO, up to 1%, had no significant effect on [<sup>3</sup>H]NMS binding under any of the experimental paradigms described above (Fig. 3).

We initially noted that there were visual decreases in cell density when cells were observed using a microscope following xanomeline pretreatments at the M1 receptor. To correct for the effects of the decrease in cell density on [<sup>3</sup>H]NMS binding we assayed protein content using the method of Bradford (Bradford, 1976) and corrected the binding curves for the changes in protein content if necessary. When cells were pretreated continuously with xanomeline in HEPES for one hour, mimicking the control treatment, there was no change in protein content at any of the receptor subtypes. Additionally, if cells were treated for one minute or one hour with xanomeline in culture media followed by washing there was no change in protein expression at any of the receptor subtypes

(Fig. 4). Thus, correction of the binding curves was not necessary following acute xanomeline treatments. When similar experiments were conducted using DMSO concentrations corresponding to those used in the preparation of xanomeline there was also no change in protein content for any of the treatment groups (Fig. 5).

### **Effects of acute carbachol treatments on inhibition of [<sup>3</sup>H]NMS binding at M1-M4 receptor subtypes**

Carbachol was employed in place of xanomeline in the pretreatment paradigm to determine if the observed effects of xanomeline pretreatments on [<sup>3</sup>H]NMS binding were unique to xanomeline or are a general property of muscarinic agonists. Carbachol was used because it is a full agonist at all of the receptor subtypes studied (Wang & El-Fakahany, 1993). There was a concentration-dependent and complete inhibition of [<sup>3</sup>H]NMS binding when carbachol was present concurrently with 0.2 nM [<sup>3</sup>H]NMS for one hour, similar to that seen with xanomeline albeit with a lower potency for all subtypes (Fig. 6; Table 1). In contrast, when cells were exposed to carbachol for one hour followed by washing and immediate use there was no significant decrease in [<sup>3</sup>H]NMS binding for any of the receptor subtypes, with the exception of cells expressing the M4 receptor, which had a concentration-dependent decrease in [<sup>3</sup>H]NMS binding (Fig. 6 d). Results are summarized in Table 1. These results are in stark contrast to those of acute wash-resistant xanomeline interaction with M1-M4 receptors (Fig. 2).

Control experiments were conducted to ascertain if pretreatment with carbachol had an effect on protein expression at any of the receptor subtypes. Cells were pretreated as described in the previous paradigm. Similar to the results presented above, acute pretreatment with carbachol had no effect on protein content for any of the subtypes (Fig. 7).



### **Effects of acute pilocarpine treatment on inhibition of [<sup>3</sup>H]NMS binding at the M3 receptor subtype**

Additional experiments were conducted at the M3 receptor to determine the effects of increasing concentrations of a partial agonist, pilocarpine, on the inhibition of [<sup>3</sup>H]NMS binding, since xanomeline is a partial agonist at the M3 receptor, whereas it is a full agonist at the M1 receptor (Chapter 4 Figs. 1 & 2). The results with pilocarpine were similar to those of carbachol at the M3 receptor. There was a concentration-dependent and complete inhibition of [<sup>3</sup>H]NMS binding when pilocarpine was present concurrently with [<sup>3</sup>H]NMS (Fig. 8; Table 1). However, when cells were treated with pilocarpine for one hour followed by washing there was no decrease in [<sup>3</sup>H]NMS binding (Fig. 8). Additionally, pretreatment with pilocarpine using the same paradigms described previously did not have an effect on protein expression for either of the treatment groups (Fig. 9).

### **The effect of acute xanomeline treatment on [<sup>3</sup>H]NMS saturation binding at M1-M4 receptor subtypes**

In subsequent experiments, cells expressing M1-M4 receptors were incubated with increasing concentrations of [<sup>3</sup>H]NMS following treatment with 10  $\mu$ M xanomeline to determine the consequence of xanomeline binding on cell-surface receptor density and radioligand-receptor affinity. In all cases, when cells were treated with 10  $\mu$ M xanomeline for one hour followed by washing and immediate use there was no significant change in the cell-surface receptor density; however, there was a significant ( $p < 0.05$ ) decrease in radioligand-receptor affinity compared to control (Fig. 10; Table 2). Noteworthy, the decrease in radioligand affinity was greatest at the M1 receptor, followed closely by the M3 receptor and then the M4 and M2 receptors (Table 2).

## **Differential effects of long-term xanomeline treatment on inhibition of [<sup>3</sup>H]NMS binding at M1-M4 receptor subtypes**

Prolonged continuous exposure to agonists has been demonstrated to lead to receptor internalization, down-regulation or uncoupling of the receptor from the G protein (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990). These effects can result in changes in the binding properties of the receptor. In subsequent experiments, we were interested in determining the long-term effects of brief and prolonged exposure to xanomeline on the binding properties across muscarinic receptor subtypes. In contrast to brief treatments followed by washing discussed above, when cells were treated with xanomeline for one minute followed by washing and incubation for 24 hours in control media a biphasic binding curve was observed at the M1 and M3 receptor subtypes, but not at the M2 or M4 receptor subtypes (Fig. 11). Complete inhibition of [<sup>3</sup>H]NMS was still observed at all receptor subtypes (Table 1). When the incubation time with xanomeline was increased to one hour prior to washing and waiting 23 hours, a biphasic binding curve was also observed at all but the M2 receptor (Fig. 11). Interestingly, there was complete inhibition of [<sup>3</sup>H]NMS binding at all but the M3 receptor under these conditions (Table 1). Similarly, when cells were incubated with xanomeline for 24 hours prior to washing a biphasic binding curve was observed at all but the M2 receptor. The high-potency site accounted for approximately 44, 41, and 29 percent of the total receptor population for the M1, M3 and M4 receptor subtypes, respectively. Interestingly, the high-potency binding site was of markedly higher potency for 24 hour xanomeline pretreatment than was observed with the acute (one minute or one hour) xanomeline pretreatments ( $p < 0.05$ ) (Table 1). Under these conditions, inhibition of [<sup>3</sup>H]NMS binding was complete at all subtypes (Table 1).

As described above for acute xanomeline treatments, control experiments using DMSO in the pretreatment paradigms were conducted to insure that the vehicle did not contribute to the observed inhibition of [<sup>3</sup>H]NMS binding. The presence of the various concentrations of DMSO used in the preparation of xanomeline had no effect on [<sup>3</sup>H]NMS binding for any of the previously described treatment paradigms (Fig. 12).

We initially noted that there were visual changes in cell density following xanomeline pretreatments at the M1 receptor. A decrease in protein content would necessitate correction of the binding curves for protein content. The method of Bradford (Bradford, 1976) was used to assess protein content and corrections to the binding curves were made as necessary. As stated above, when cells were pretreated continuously with xanomeline in HEPES buffer for one hour (control) there was no change in protein content at any of the receptor subtypes (Fig. 4). However, when cells expressing the M1 receptor were treated with xanomeline for one minute or one hour followed by washing and overnight incubation in ligand free media there was a concentration dependent decrease in protein content (Fig. 13 a). Similarly, 24 hour-treatment followed by washing also resulted in a decrease in protein content at the M1 receptor. Thus, the binding curves at the M1 receptor were corrected as a result of xanomeline pretreatment. In contrast, when the same pretreatments were performed at the M2-M4 receptors there was no change in protein content for any of the treatment paradigms (Fig. 13 b-d). When similar experiments were conducted utilizing the same concentrations of DMSO used in the preparation of xanomeline there was no change in protein content for any of the pretreatment paradigms described previously at the various receptor subtypes (Fig. 14).

## **Effects of long-term carbachol treatments on inhibition of [<sup>3</sup>H]NMS binding at the M1-M4 receptor subtypes**

To ascertain whether the long-term effects of xanomeline binding were specific to xanomeline or are a general property of muscarinic agonist binding, experiments were undertaken using the full agonist carbachol. When cells were exposed to increasing concentrations of carbachol for one hour followed by washing and further incubation for 23 hours in control media there was no decrease in [<sup>3</sup>H]NMS binding for any of the receptor subtypes (Fig. 15). This is in stark contrast to the delayed appearance of biphasic binding curves at all but the M2 receptor subtype with xanomeline treatment (Fig. 11). However, when cells were exposed to carbachol for 24 hours prior to washing there was a concentration-dependent decrease in [<sup>3</sup>H]NMS binding at all of the subtypes (Fig. 15). The maximal decrease in binding was significantly ( $p < 0.05$ ) greater for the M1 and M2 receptors in comparison to the M3 and M4 receptors. The results are summarized in Table 1. Again, this was in stark contrast to the effects of 24 hour-treatment with xanomeline prior to washing that resulted in a biphasic binding curve at all except the M2 receptor and complete inhibition of [<sup>3</sup>H]NMS binding at all subtypes (Fig. 11). These data indicate that xanomeline binding results in long-term effects that are distinct from the conventional reversible agonist carbachol. To determine if there were changes in protein content following any of the long-term exposure to carbachol paradigms protein assays using the method of Bradford (Bradford, 1976) were conducted. There was a decrease in protein content at the M1 receptor following 24 hour carbachol treatment, thus necessitating corrections be made to the binding curve (Fig. 16 a). However, there was no change in protein for any of the other previously described carbachol pretreatments across receptor subtypes (Fig. 16).

### **Effects of long-term pilocarpine on inhibition of [<sup>3</sup>H]NMS binding at the M3 receptor subtype**

Additional experiments were conducted at the M3 receptor using pilocarpine in the inhibition of [<sup>3</sup>H]NMS binding assay instead of carbachol to further determine if the long-term effects of exposure to a conventional agonist are different from xanomeline. As noted previously, pilocarpine is a partial agonist at the M3 receptor. When cells were exposed to increasing concentrations of pilocarpine for one hour followed by washing and waiting 23 hours there was no decrease in [<sup>3</sup>H]NMS binding (Fig. 17). However, when cells were pretreated with pilocarpine for 24 hours prior to washing there was a concentration-dependent decrease in [<sup>3</sup>H]NMS binding similar to that observed with carbachol, however the potency was lower for pilocarpine compared to carbachol (Fig. 17). Results are summarized in Table 1. Taken together, these data demonstrate that the long-term effects of xanomeline binding are distinct from those of conventional muscarinic agonists. When experiments were conducted to determine protein content following pilocarpine treatment paradigms as discussed above, no differences were seen between any of the pretreatment groups (Fig. 18).

### **The effects of long-term xanomeline treatment on [<sup>3</sup>H]NMS saturation binding at M1-M4 receptor subtypes**

Experiments were conducted to determine the long-term effects of 10 μM xanomeline exposure on cell-surface receptor density and radioligand-receptor affinity. When cells were treated with xanomeline for one hour followed by washing and incubation for 23 hours in the absence of free ligand there was no change in radioligand-receptor affinity at any of the subtypes, but a significant ( $p < 0.05$ ) decrease in cell-surface receptor density at all except the M2 receptor (Fig. 19). When cells were treated

with xanomeline for 24 hours prior to washing there was a significant ( $p < 0.05$ ) decrease in radioligand-receptor affinity across all four subtypes and a significant ( $p < 0.05$ ) decrease in cell-surface receptor density at all except the M2 subtype (Fig. 19). Interestingly, the decrease in cell-surface receptor density was more pronounced at the M1 receptor compared to the M3 or M4 receptors. Results are summarized in Table 2.

### **The effects of long-term conventional agonist treatment on [<sup>3</sup>H]NMS saturation binding parameters at the M3 receptor**

Additional experiments were conducted to determine if the effects of prolonged exposure of cells to the conventional agonists carbachol or pilocarpine are distinct from those of xanomeline. Previous research has shown that prolonged exposure of cells expressing muscarinic receptors to carbachol leads to a decrease in cell-surface receptor density with no change in radioligand affinity (Cioffi & El-Fakahany, 1989; Hu et al., 1991; May et al., 2005; van de Westerlo et al., 1995; Wang et al., 1990; Yang et al., 1993). To confirm these results in our experimental paradigm, we determined the effects of long-term exposure of M3 receptors to carbachol. CHO cells expressing the M3 receptor were subjected to the various pretreatments and washing with 10  $\mu$ M carbachol, then exposed to increasing concentrations of [<sup>3</sup>H]NMS in order to determine changes in cell-surface receptor density and radioligand affinity. When cells were treated with carbachol for one hour followed by washing and incubation in control media for 23 hours, there was no change in the maximal binding or radioligand affinity compared to control cells (Fig. 20 a). In contrast, pretreatment with xanomeline under the same conditions resulted in a decrease in cell-surface receptor density (Fig. 19). In agreement with previous literature (Hu et al., 1991; May et al., 2005; van de Westerlo et al., 1995; Yang et al., 1993), pretreatment with carbachol for 24 hours prior to washing

resulted in a decrease in maximal binding. However, there was a significant ( $p < 0.05$ ) increase in radioligand affinity (Fig. 20 a; Table 2). This is in stark contrast to the marked decrease in radioligand affinity observed following pretreatment with xanomeline for 24 hours (Fig. 19).

Previous research regarding the effects of prolonged exposure to pilocarpine has produced mixed findings. Shifrin & Klein (1980) found that prolonged exposure to pilocarpine resulted in a decrease in receptor density. In contrast, another report suggested that pilocarpine has little effect on the actual number of receptors, as the apparent down-regulation was due to pilocarpine being released when cells were lysed prior to the binding assay (Milligan & Strange, 1983). In our paradigm, when cells were exposed to 10  $\mu\text{M}$  pilocarpine for one hour followed by washing and waiting 23 hours there was a slight, but significant ( $p < 0.05$ ), decrease in cell surface receptor density without a change in receptor affinity (Fig. 20 b). Additionally in our paradigm, 24 hour-exposure of cells to pilocarpine prior to washing resulted in a significant ( $p < 0.05$ ) decrease in receptor density along with a significant ( $p < 0.05$ ) increase in radioligand affinity (Fig. 20 b). Results are summarized in Table 2. These results further demonstrate a unique binding profile following prolonged exposure to xanomeline compared to conventional agonists.

## **Discussion**

The high degree of sequence homology of the orthosteric binding site of various subtypes of muscarinic receptors has made it difficult to develop an agonist that binds with selectivity at a particular subtype (Wess, 2004). Xanomeline is thought to be an M1/M4 functionally selective agonist, although it can bind at all subtypes (Bymaster et al., 1997; Shannon et al., 1994). Xanomeline has been shown to bind in a unique

manner, with both reversible and wash-resistant binding at the M1 receptor (Christopoulos et al., 1998). Thus, experiments were undertaken to determine the short- and long-term effects of xanomeline binding at the M1-M4 muscarinic receptor subtypes expressed in CHO cells.

Our results indicate that wash-resistant binding occurs rapidly (within 15 seconds) at all four receptor subtypes studied (Fig. 1). However, the immediate decrease in [<sup>3</sup>H]NMS binding was greater at the M1 and M3 subtypes compared to the M2 and M4 subtypes, as was the maximal inhibition of [<sup>3</sup>H]NMS binding. It is possible that the differences in the speed of binding are a result of differences in the structure of the allosteric site on receptor subtypes. Alternatively, it might be due to differences in receptor conformation induced by receptor subtype coupling to different G proteins and the subsequent activation of different signaling pathways. Both the M1 and M3 receptors couple to G<sub>q/11</sub> G proteins (Volpicelli & Levey, 2004) whereas the M2 and M4 receptors couple to G<sub>i/o</sub> G proteins (Eglen, 2005).

In agreement with previous research (Bymaster et al., 1997; Shannon et al., 1994) our results demonstrate that xanomeline is able to bind to all muscarinic receptor subtypes. This can be seen as a concentration-dependent decrease in [<sup>3</sup>H]NMS binding at all four receptor subtypes when xanomeline is present concurrently with the radioligand in the binding assay (Fig 2). In addition, when cells were incubated with xanomeline for one minute or one hour prior to washing and immediately used in the binding assay there was a concentration-dependent decrease in [<sup>3</sup>H]NMS binding, demonstrating the ability of xanomeline to bind in a persistent manner at all subtypes. This is in agreement with previously published data that wash-resistant binding occurs at the M1 (Christopoulos & El-Fakahany, 1997), and M2 (Jakubik et al., 2002; Jakubik et al., 2006) receptors. Interestingly, the rank order of potencies at various receptor



subtypes is different between the various xanomeline treatments. When cells are treated concurrently with xanomeline and [<sup>3</sup>H]NMS the rank order is M3 ≈ M4 > M2 > M1. This protocol detects both xanomeline reversible and wash-resistant binding. The reverse rank order of potency (M1 > M2 > M3 ≈ M4) was observed when cells are pretreated with xanomeline for one minute followed by washing and immediate use. This might be explained, at least in part, by different kinetics of the rapid phase of xanomeline binding at the different subtypes, especially since differences in potency among subtypes disappear when cells are treated with xanomeline for one hour followed by washing and immediate determination of radioligand binding.

When parallel experiments were undertaken utilizing carbachol in place of xanomeline in the various pretreatment paradigms major differences were observed in the binding profiles of the two agonists (Fig 6). There was a concentration-dependent decrease in [<sup>3</sup>H]NMS binding with carbachol when concurrently present with the radioligand at all subtypes, similar to that seen with xanomeline; however, the potency was markedly decreased. Carbachol pretreatment for one hour followed by washing and immediate use did not result in a decrease in [<sup>3</sup>H]NMS binding for any of the subtypes, except when cells expressing the M4 receptor were treated for one hour followed by washing and immediate use. This was in stark contrast to the monophasic binding curves observed with acute xanomeline treatments. The decrease in [<sup>3</sup>H]NMS binding observed at the M4 receptor following acute carbachol treatment could be due to rapid internalization of the receptor (Feigenbaum & El-Fakahany, 1984; Holroyd et al., 1999). Additionally, pretreatment with pilocarpine at the M3 receptor resulted in a binding profile similar to that observed with carbachol (Fig. 8). Together these results demonstrate that xanomeline interacts with muscarinic receptors in a manner that is distinct from conventional agonists.

Xanomeline was shown to bind in a wash-resistant manner at all four receptor subtypes. Thus, additional experiments were undertaken to determine if the acute wash-resistant binding of xanomeline leads to subtype-dependent changes in receptor density or radioligand affinity. Interestingly, there was no change in receptor density following acute xanomeline treatment compared to control at any of the receptor subtypes (Fig. 10). Similarly, all receptor subtypes demonstrated a decrease in radioligand affinity following acute treatment with xanomeline, albeit to slightly different degrees. Taken together, these data demonstrate that the effects of acute xanomeline treatment are similar across muscarinic receptor subtypes.

Regulation of receptors is time dependent, with receptor internalization taking place at shorter durations of exposure to agonists, and down-regulation predominating at longer incubation times (Maloteaux & Hermans, 1994). In addition, allosteric modulators can modify the conformation of the orthosteric site resulting in changes in the binding properties of additional agonists or antagonists (Christopoulos, 2002). Thus, the effects of prolonged treatment with xanomeline were investigated across the four receptor subtypes. Treatment of cells with xanomeline for one minute followed by washing and incubation in control media for 24 hours results in a monophasic binding curve for the M2 and M4 receptors, but a biphasic binding curve at the M1 and M3 receptors (Fig. 11). However, increasing the pretreatment time to one hour prior to washing and waiting results in biphasic binding curves at all but the M2 receptor (Fig. 11). The observed differences in the long-term effects of exposure of M4 receptors to xanomeline for one minute or one hour suggest a time-dependent interaction of xanomeline with different receptor conformations. Long-term treatment with agonists has been shown to result in internalization/down-regulation of receptors (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990). Thus, the development of the apparent high-potency site at the M1, M3 and M4

receptors could be a reflection of internalization/down-regulation of the receptor that results in a decrease in radioligand binding to the receptor. In support of this notion, the effects of one hour pretreatment with xanomeline followed by washing and waiting were mimicked by treatment with xanomeline for 24 hours. However, the latter condition resulted in higher potencies at both the high and low potency sites.

Parallel experiments utilizing carbachol in place of xanomeline in the binding paradigms described above further demonstrated that xanomeline interacts with the muscarinic receptors in a unique manner. Carbachol pretreatment for one hour followed by washing and further incubation for 23 hours in control media did not result in a decrease in [<sup>3</sup>H]NMS binding for any of the subtypes. When cells were treated with carbachol for 24 hours followed by washing there was a concentration-dependent decrease in [<sup>3</sup>H]NMS binding at all of the subtypes, however, the resultant decrease was best-fit by a monophasic binding curve (Fig. 15). The results of either of these treatments were in stark contrast to the biphasic binding curve observed with xanomeline pretreatments under the same conditions. Interestingly, exposure of cells expressing the M3 receptor to pilocarpine using the various pretreatment paradigms results in a binding profile similar to that seen with carbachol, albeit with different potencies (Fig. 17). Taken together these results further demonstrate that long-term exposure to xanomeline results in a ligand-receptor interaction that is distinct from that induced by conventional muscarinic receptor agonists. Additionally, receptor modulation of the M2 receptor by xanomeline is distinctly different from the other receptor subtypes. The results following the exposure of M2 receptors to carbachol demonstrate that the lack of similar long-term effects of xanomeline at the M2 receptor compared to the other subtypes is not due to insensitivity of this receptor to agonist-induced regulation, since the effects of carbachol are similar to those seen at other subtypes. We propose that

the lack of long-term effects of xanomeline at the M2 receptor may be due to absence or a very minimal functional response to xanomeline. It is also important to consider that since xanomeline is able to bind, but stimulates a minimal if any functional response, it may have antagonistic properties at the M2 receptor.

To further support the idea that the effects of xanomeline may be a result of allosteric receptor modulation, the ability of xanomeline to inhibit [<sup>3</sup>H]NMS binding was studied. Incomplete inhibition of radioligand binding is a hallmark of allosteric receptor modulation (Lee & El-Fakahany, 1991). When cells were treated with xanomeline for one minute or one hour followed by washing and waiting there was incomplete inhibition of [<sup>3</sup>H]NMS binding at all subtypes, except for one hour treatment at the M3 receptor (Fig. 11). Thus, it is possible that the long-term effects of xanomeline binding are due to changes in cooperativity between the orthosteric and allosteric binding sites. However, it is also possible that the lack of complete inhibition of [<sup>3</sup>H]NMS binding is a result of the inability to test higher concentrations of xanomeline. Interestingly, when cells were treated with xanomeline for 24 hours prior to washing there was once again complete inhibition of [<sup>3</sup>H]NMS binding suggesting receptor modulation through internalization/down-regulation (Table 1).

Further experiments were undertaken to quantitate the effects of xanomeline binding on cell-surface receptor density and radioligand affinity (Fig. 19). When cells expressing M1, M3 or M4 receptors were exposed to xanomeline for one hour followed by washing and 23 hour incubation in ligand-free media or treated with xanomeline for 24 hours prior to washing there was a decrease in cell-surface receptor density. However, the decrease was more distinctive at the M1 receptor compared to the other subtypes. These results are in agreement with the results of experiments of inhibition of [<sup>3</sup>H]NMS binding in which the M1 receptor had a more distinctive high-potency binding

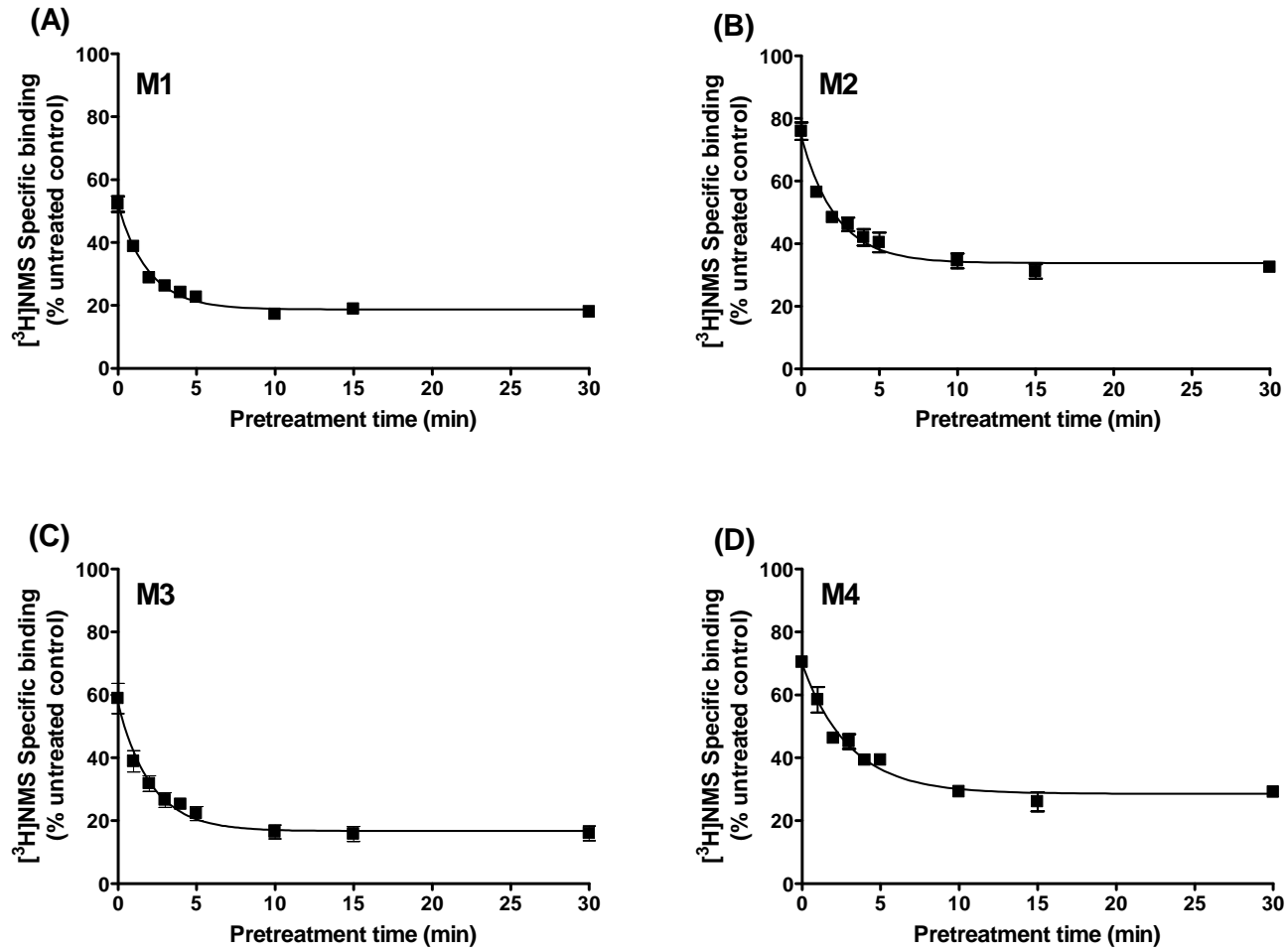
site compared to the M3 or M4 receptors (Fig. 11). Prolonged exposure to agonists has been shown to lead to receptor internalization and/or down-regulation (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990). This provides further support to the notion that the decrease in cell-surface receptor density following long-term xanomeline exposure could be a result of internalization or down-regulation of the receptor. In contrast, neither xanomeline pretreatment condition had an effect on cell-surface receptor density in cells expressing the M2 receptor. This further demonstrates that the long-term regulation of the M2 receptor by xanomeline is distinct from the other receptor subtypes and suggests the possibility that xanomeline is unable to activate the M2 receptor. In addition, it should be noted that the cell-surface receptor density levels in our experimental system are similar for the M1, M3 and M4 receptors, but are considerably lower at the M2 receptor (Table 2). The decreased cell surface receptor density at the M2 receptor is not likely to lead to the large differences in the effect of long-term xanomeline exposure compared to the other receptor subtypes. These differences are more likely to be the result of varied efficacy of xanomeline at various receptor subtypes. Previous work by Shannon et al. (1994) found that xanomeline had little efficacy, if any, in heart tissue that predominantly has M2 receptors. Additionally, recent results quantifying ERK phosphorylation showed that xanomeline only elicited 40% of the functional response at the M2 receptor compared to the M4 receptor in cells that had similar receptor expression (Arthur Christopoulos, personal communication). In contrast to effects on receptor expression, radioligand affinity was decreased following 24 hour xanomeline treatment prior to washing at all four subtypes. This suggests that the change in radioligand affinity is dependent solely on binding of xanomeline to the receptor.

Parallel experiments were conducted at the M3 receptor using carbachol or pilocarpine in place of xanomeline in the pretreatment paradigms. Previous research

has demonstrated that exposure of cells to carbachol for 24 hours results in a decrease in cell-surface receptor density (Hu et al., 1991; May et al., 2005; van de Westerlo et al., 1995; Yang et al., 1993). In either case, pretreatment for one hour followed by washing and waiting resulted in no change in radioligand affinity and only a slight decrease in receptor density following pilocarpine treatment. Twenty-four hour treatment with either carbachol or pilocarpine resulted in a decrease in cell-surface receptor density; however, this decrease was less marked than the decrease observed with the xanomeline treatment following the same experimental protocol. Additionally, 24-hour treatment with either agonist resulted in an increase in radioligand affinity, which is in stark contrast to the decrease in radioligand affinity observed with xanomeline treatment. Together, these results further demonstrate that xanomeline interacts in a unique manner with muscarinic receptors.

In summary, we have shown that although xanomeline can bind in both a reversible and wash-resistant manner with near equal potency across the M1-M4 receptor subtypes, the long-term effects of xanomeline binding are distinctively different at the M2 receptor compared to the other receptor subtypes. The differences in the long-term effects of xanomeline are most likely due to a difference in xanomeline efficacy between subtypes. Additionally, the effects of xanomeline treatment are distinctively different from the effects of treatment with conventional muscarinic receptors employing the same paradigms. Studies described in subsequent chapters are aimed at determining the effects of xanomeline on the functional properties of the M1 and M3 receptors and the mechanisms of receptor regulation by xanomeline.

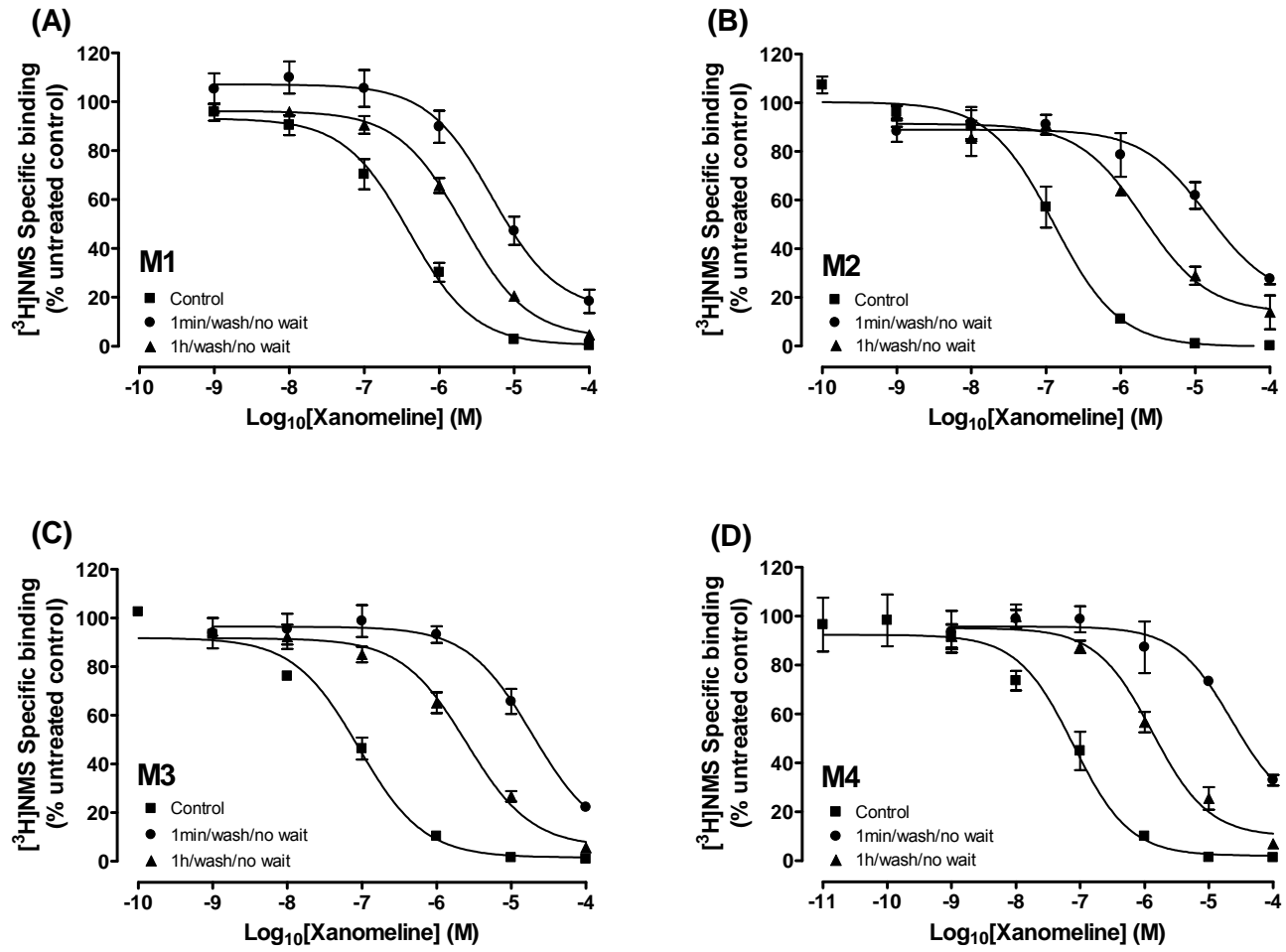
Figure 1



**Figure 1. Time course of the development of xanomeline wash-resistant binding at hM1-hM4 muscarinic receptor subtypes expressed in CHO cells.** Cells were treated with 10  $\mu$ M xanomeline for increasing lengths of time (< 15 seconds to 30 minutes) followed by washing and incubation with 0.2 nM [ $^3$ H]NMS for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Binding in the absence of xanomeline pretreatment is denoted by 100%. Individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 7400, 7600, 5700 and 9900 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 3 experiments conducted in triplicate.

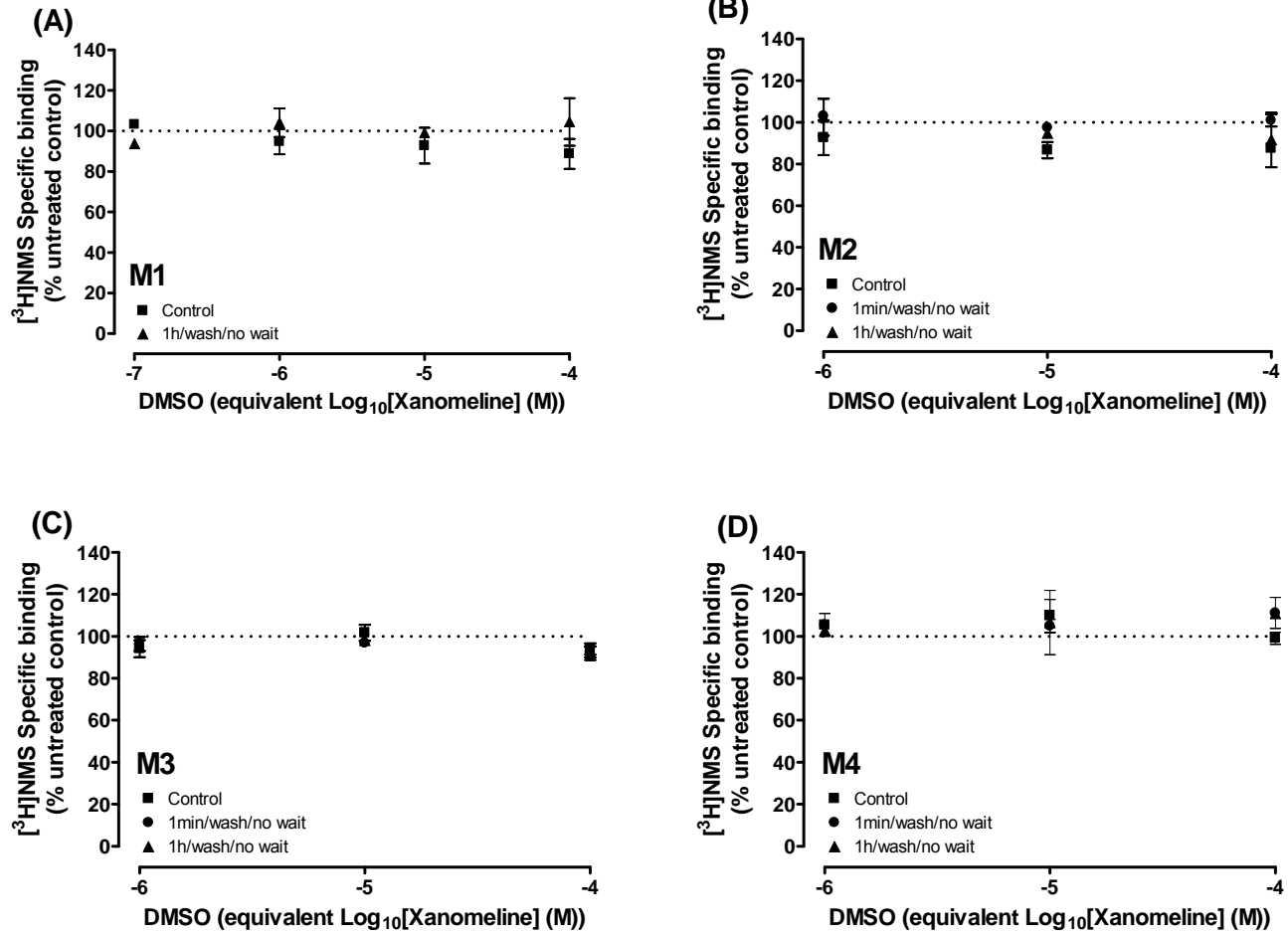


Figure 2



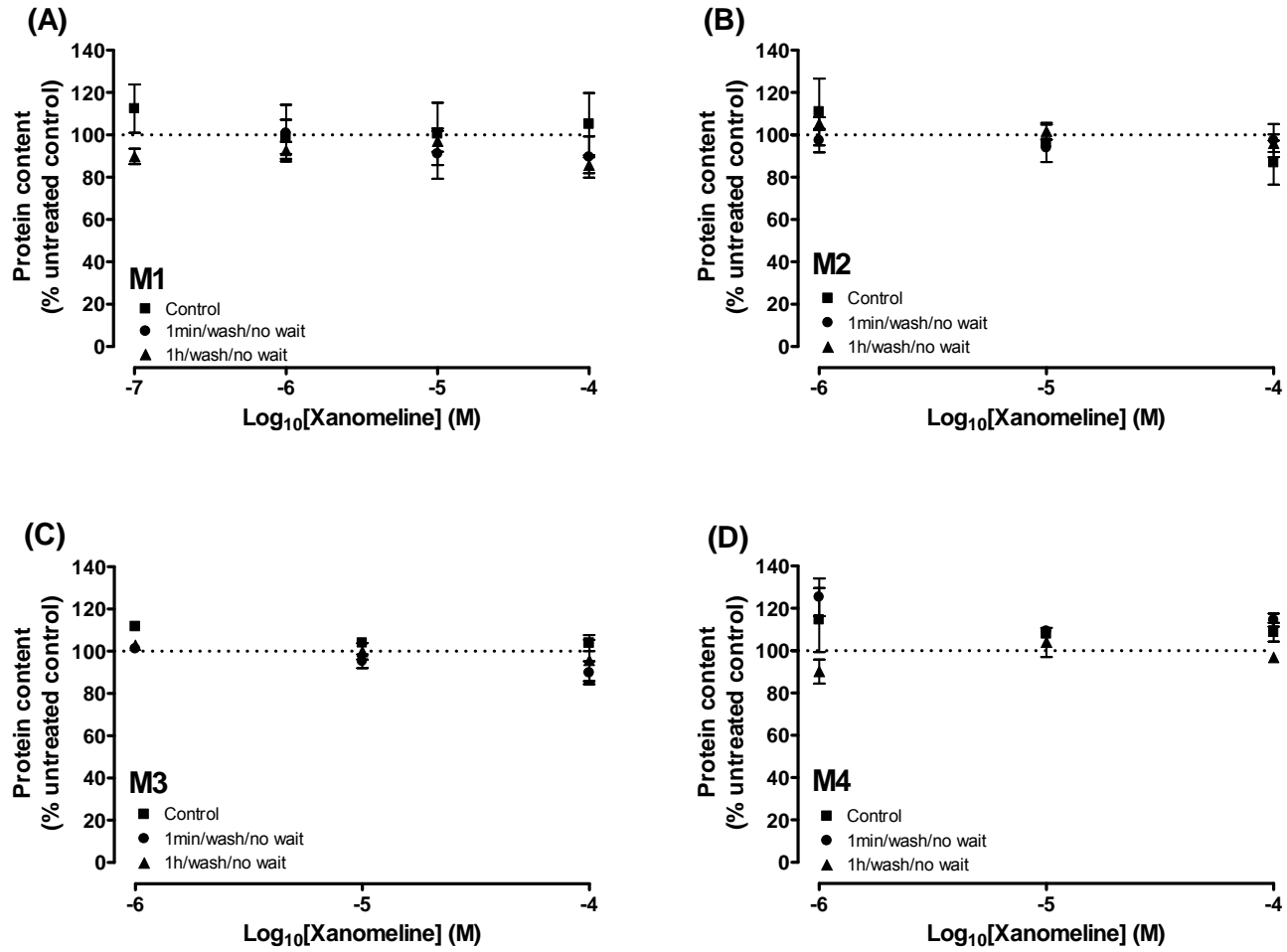
**Figure 2. Inhibition of [<sup>3</sup>H]NMS binding for acute xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of xanomeline for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing. Subsequently, cells were incubated for 1 hour at 37°C with 0.2 nM [<sup>3</sup>H]NMS concurrently with xanomeline (closed squares) or following xanomeline pretreatments. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 13200, 12100, 21300 and 19000 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 3-9 experiments conducted in triplicate.

Figure 3



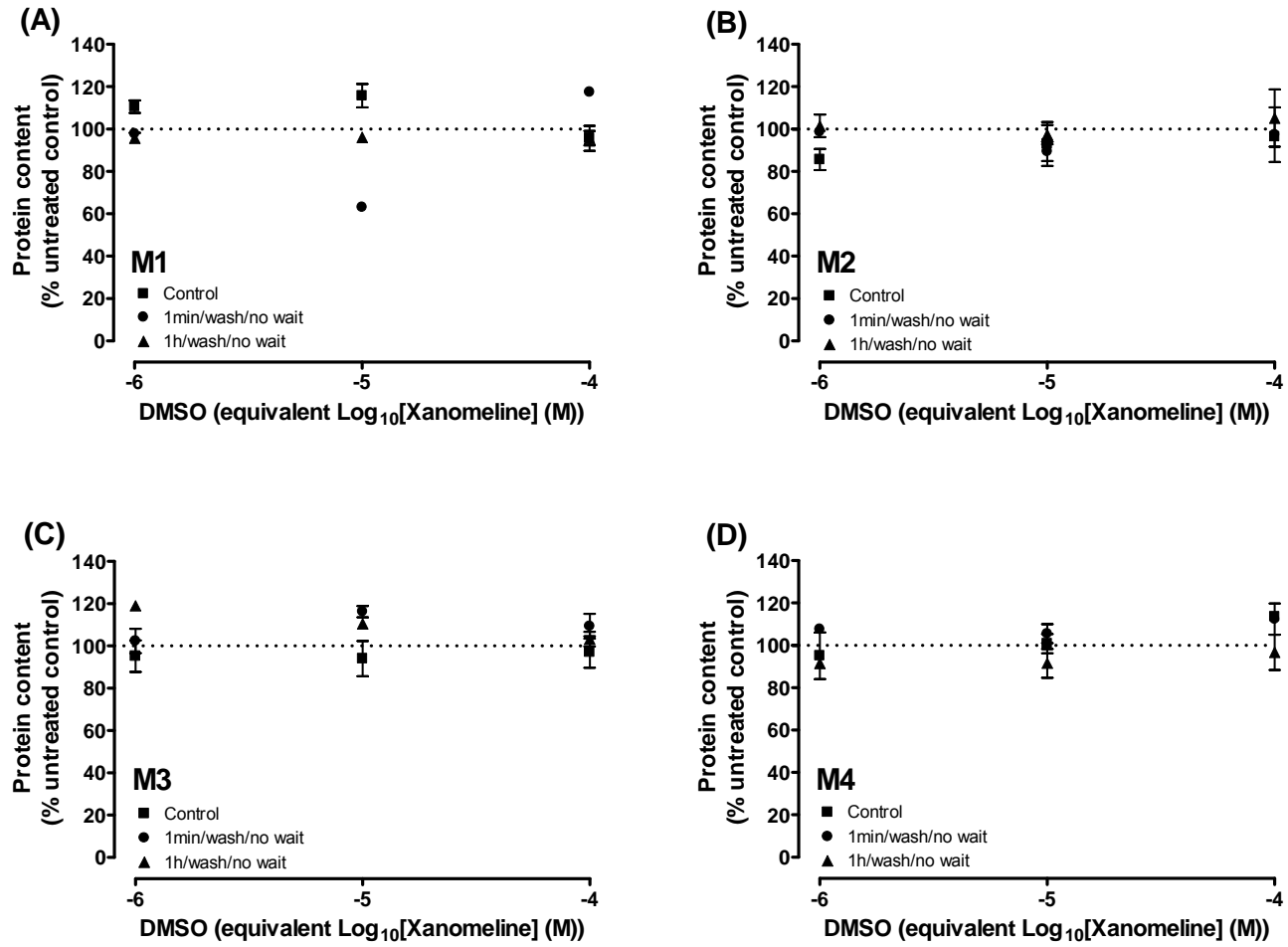
**Figure 3. Effects of acute DMSO pretreatments on the inhibition of [<sup>3</sup>H]NMS binding in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of DMSO in media (corresponding to the concentrations used in the preparation of xanomeline) for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing. Subsequently, cells were exposed concurrently to DMSO and 0.2 nM [<sup>3</sup>H]NMS (closed squares) or to [<sup>3</sup>H]NMS following DMSO pretreatments for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control binding in the absence of DMSO treatment which had a mean of 11900, 15800, 21300 and 15400 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 2 experiments conducted in triplicate. Data from (A) provided by Marianne Grant.

Figure 4



**Figure 4. Effects of acute xanomeline pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of xanomeline for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing. Subsequently, reactions were terminated using 1 M NaOH and protein content on untreated cells incubated with xanomeline in HEPES buffer for 1 hour (closed squares) or xanomeline pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of xanomeline treatment which had a mean of 130, 256, 153 and 196  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2-3 experiments conducted in triplicate.

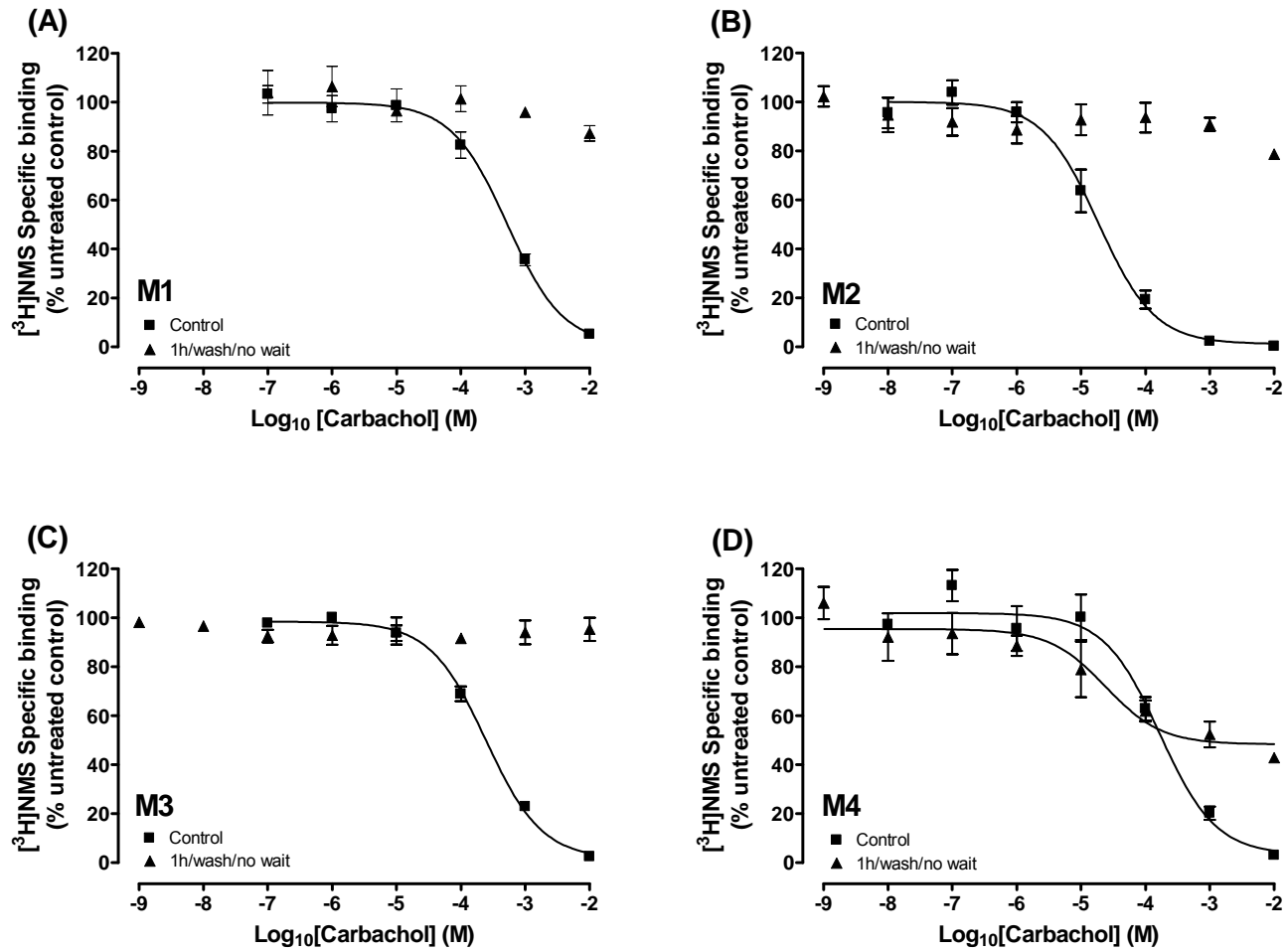
Figure 5



**Figure 5. Effects of acute DMSO pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of DMSO in media (corresponding to the concentrations used in the preparation of xanomeline) for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing. Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with DMSO in HEPES buffer for 1 hour (closed squares) or DMSO pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of DMSO treatment which had a mean of 130, 265, 180 and 196  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

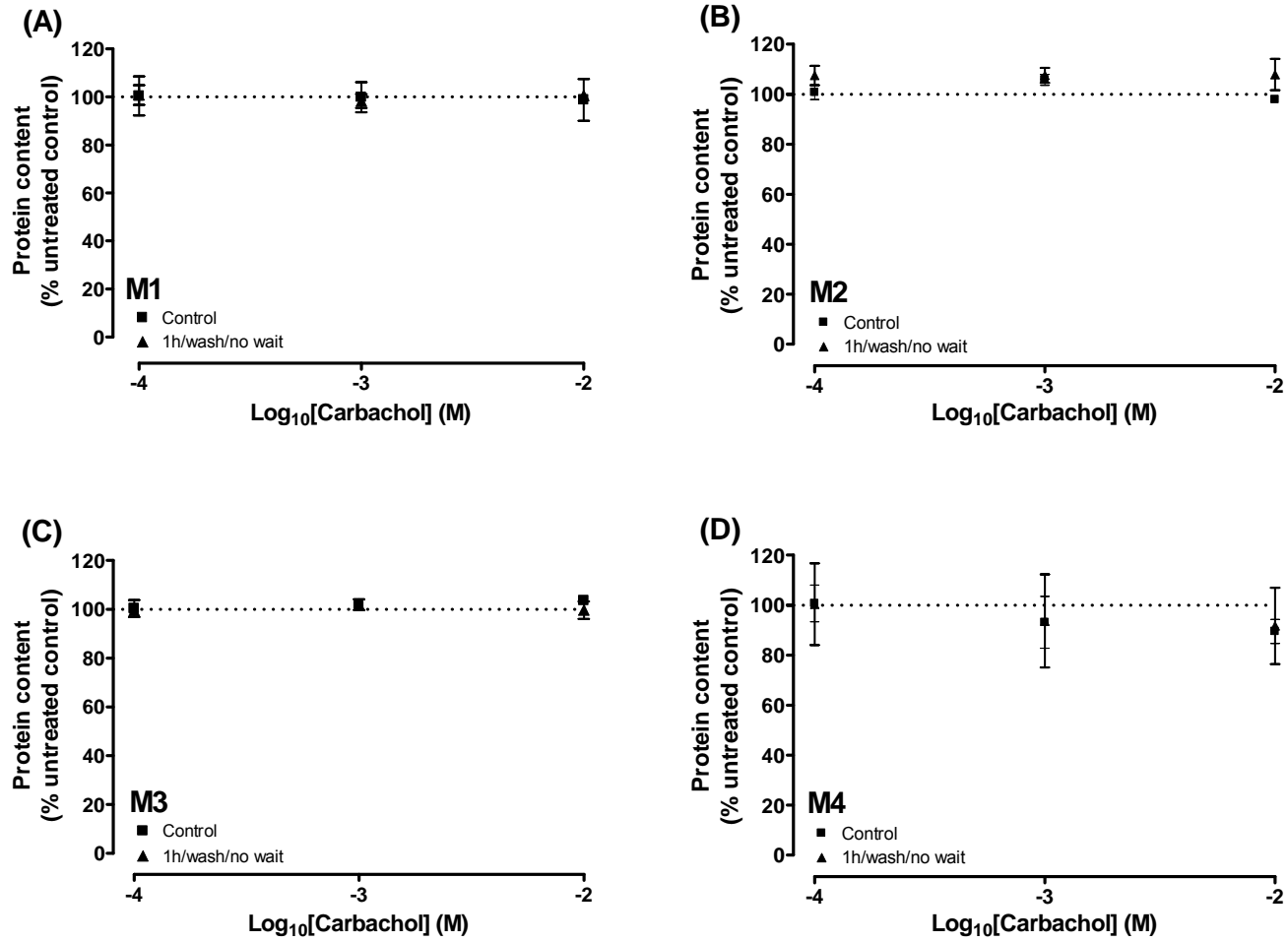


Figure 6



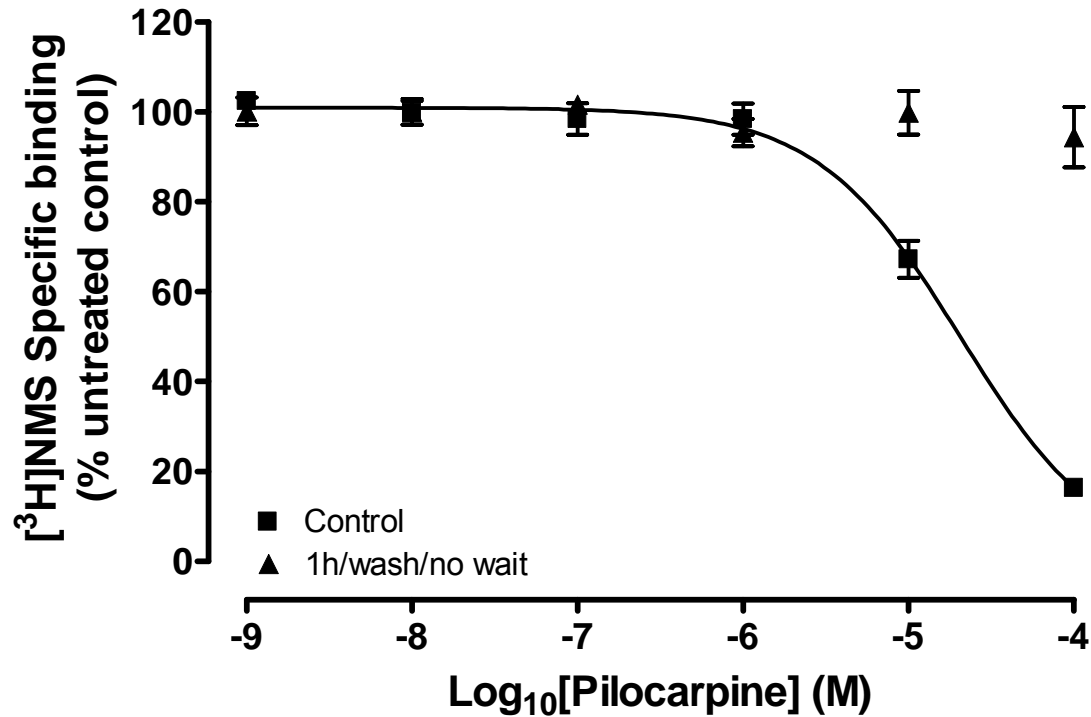
**Figure 6. Inhibition of [<sup>3</sup>H]NMS binding for acute carbachol pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of carbachol for 1 hour followed by washing (closed triangles). Subsequently, cells were incubated with 0.2 nM [<sup>3</sup>H]NMS concurrently with carbachol (closed squares) or following carbachol pretreatments for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Individual experiments were normalized to control binding in the absence of carbachol treatment which had a mean of 13000, 16500, 17700, and 16100 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 3-4 experiments conducted in triplicate.

Figure 7



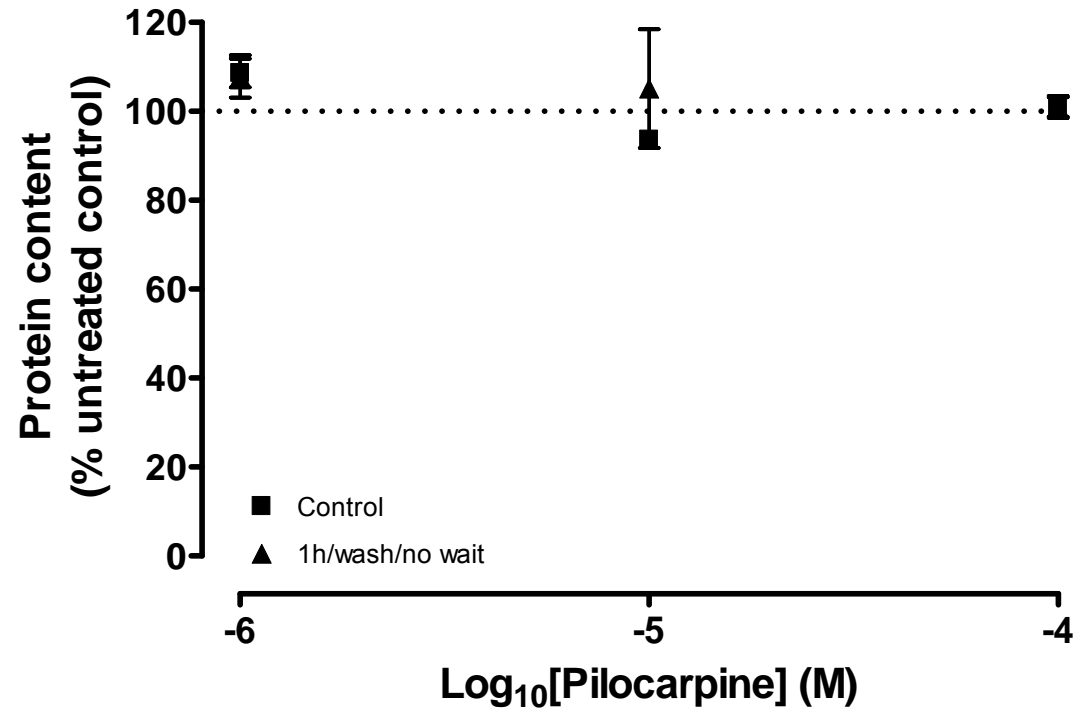
**Figure 7. Effects of acute carbachol pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of carbachol for 1 hour (closed triangles) followed by washing. Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with carbachol in HEPES buffer for 1 hour (closed squares) or carbachol pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of carbachol treatment which had a mean of 180, 270, 153 and 296  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

Figure 8



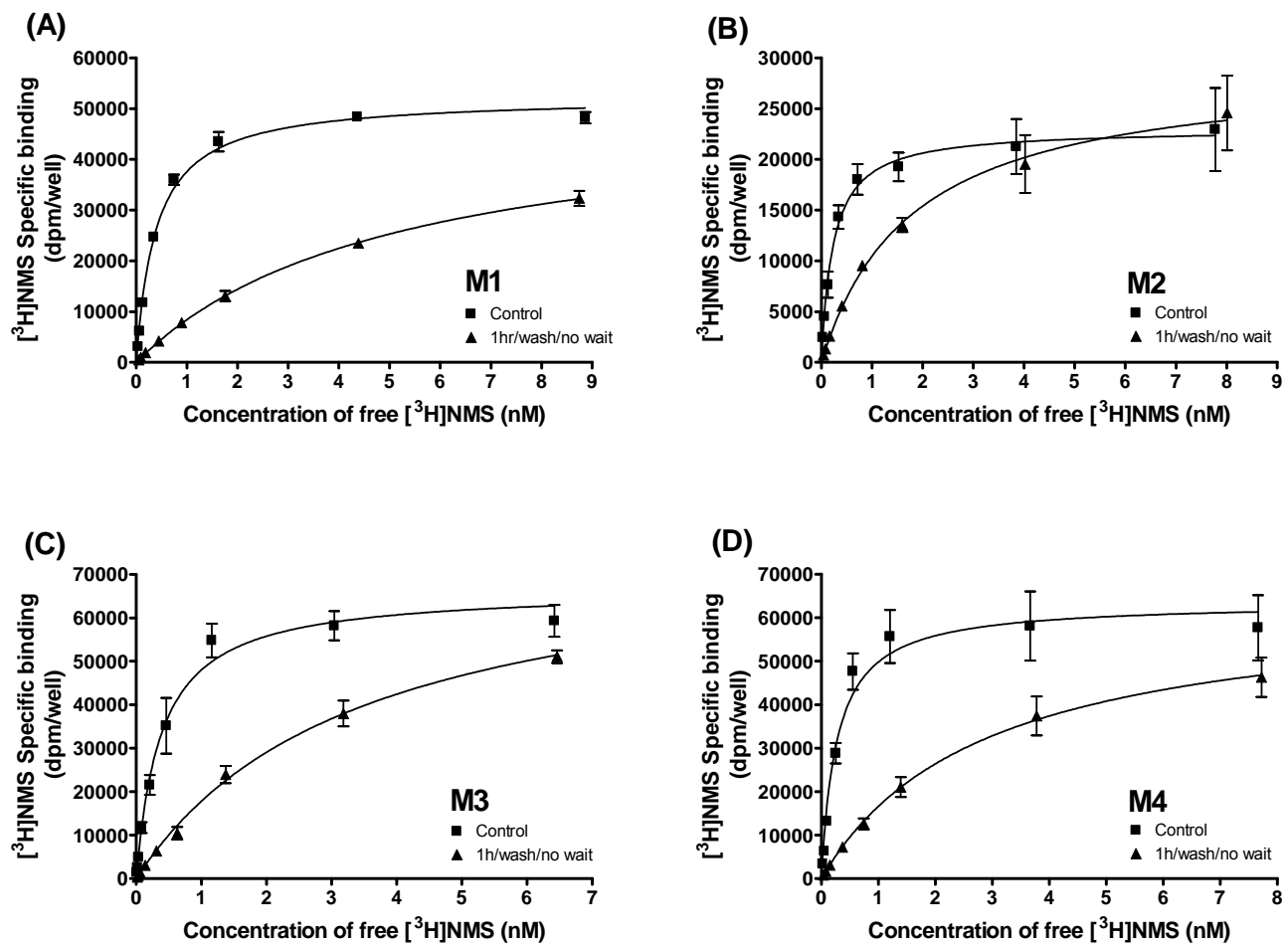
**Figure 8. Inhibition of  $[^3\text{H}]\text{NMS}$  binding for acute pilocarpine pretreatments in CHO cells stably expressing the hM3 receptor subtype.** Cells were pretreated with increasing concentrations of pilocarpine for 1 hour followed by washing (closed triangles). Subsequently, cells were incubated with 0.2 nM  $[^3\text{H}]\text{NMS}$  concurrently with pilocarpine (closed squares) or following pilocarpine pretreatments for 1 hour at 37°C. Individual experiments were normalized to control binding in the absence of pilocarpine treatment which had a mean of 17500 dpm/well. Values represent the means  $\pm$  S.E. of 3 experiments conducted in triplicate.

Figure 9



**Figure 9. Effects of acute pilocarpine pretreatments on protein content in CHO cells stably expressing the hM3 receptor subtype.** Cells were pretreated with increasing concentrations of pilocarpine for 1 hour (closed triangles) followed by washing. Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with pilocarpine in HEPES buffer for 1 hour (closed squares) or pilocarpine pretreated cells was determined using the method of Bradford (Bradford, 1976). Individual experiments were normalized to control protein in the absence of pilocarpine treatment which had a mean of 180  $\mu$ g/well. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

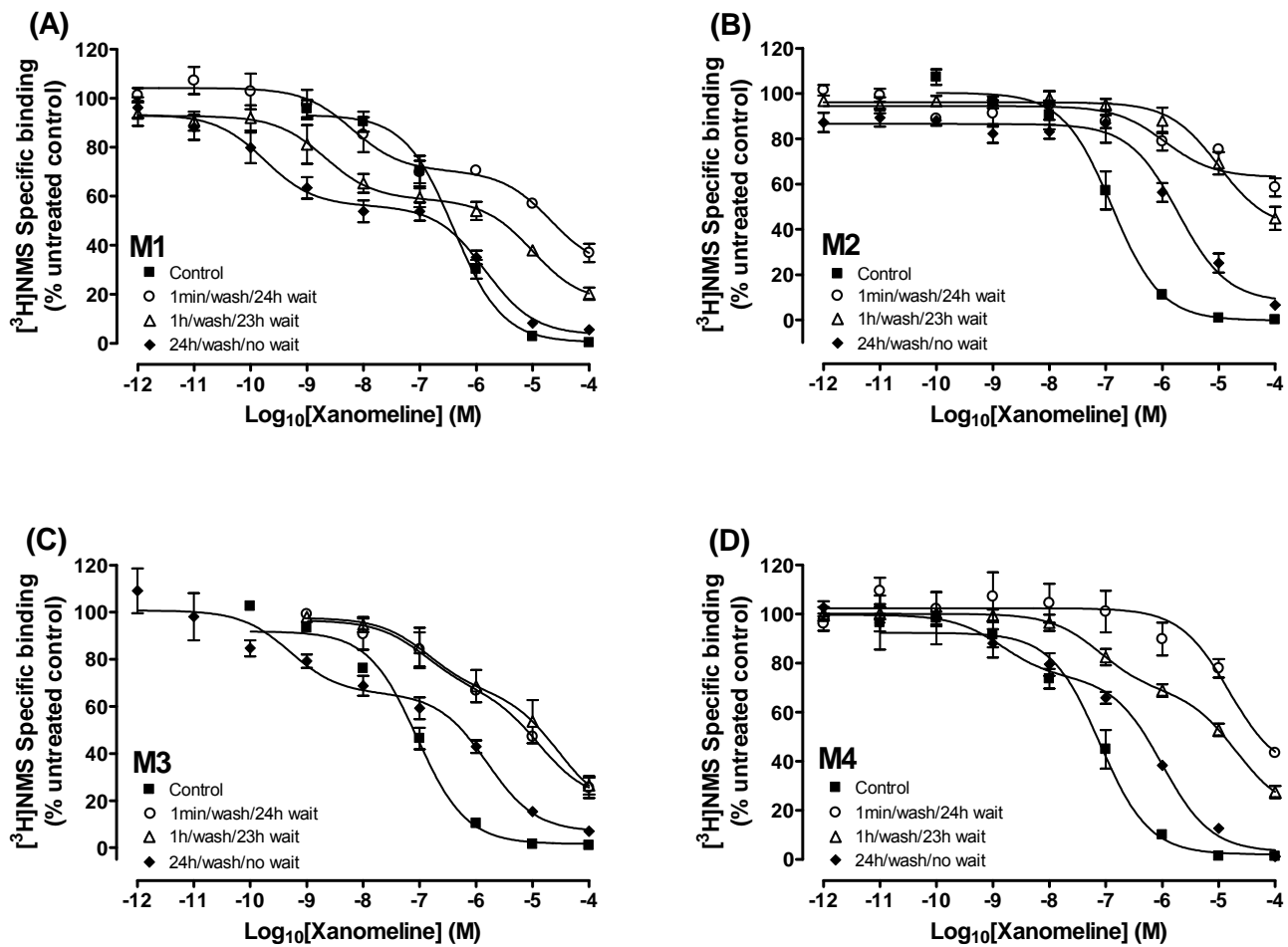
Figure 10



**Figure 10. [<sup>3</sup>H]NMS saturation binding subsequent to acute xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with 10 μM xanomeline for 1 hour followed by washing (closed triangles). Subsequently, untreated (closed squares) or xanomeline pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 8.9 nM) for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Values represent the means ± S.E. of 3-4 experiments conducted in triplicate.

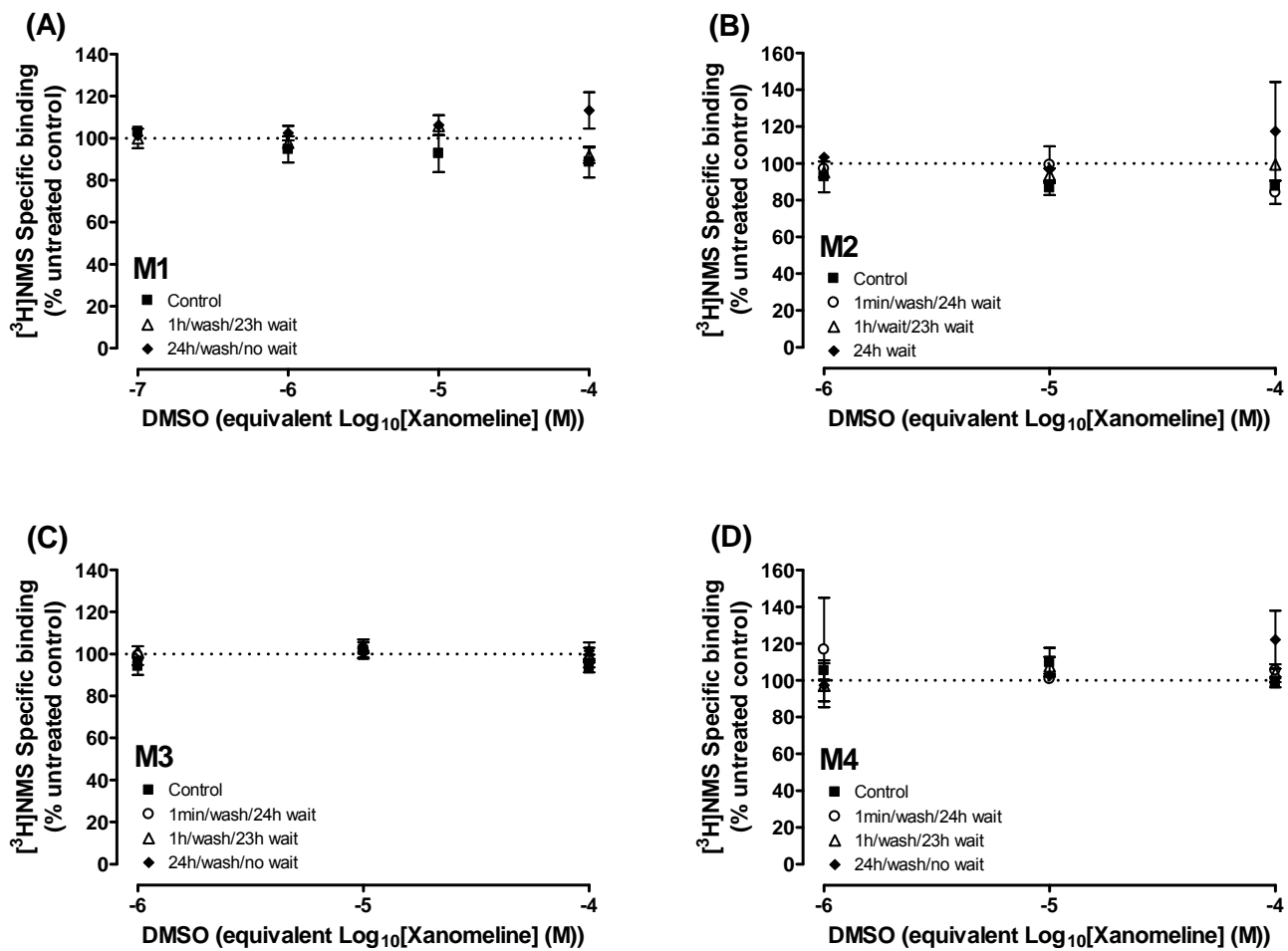


Figure 11



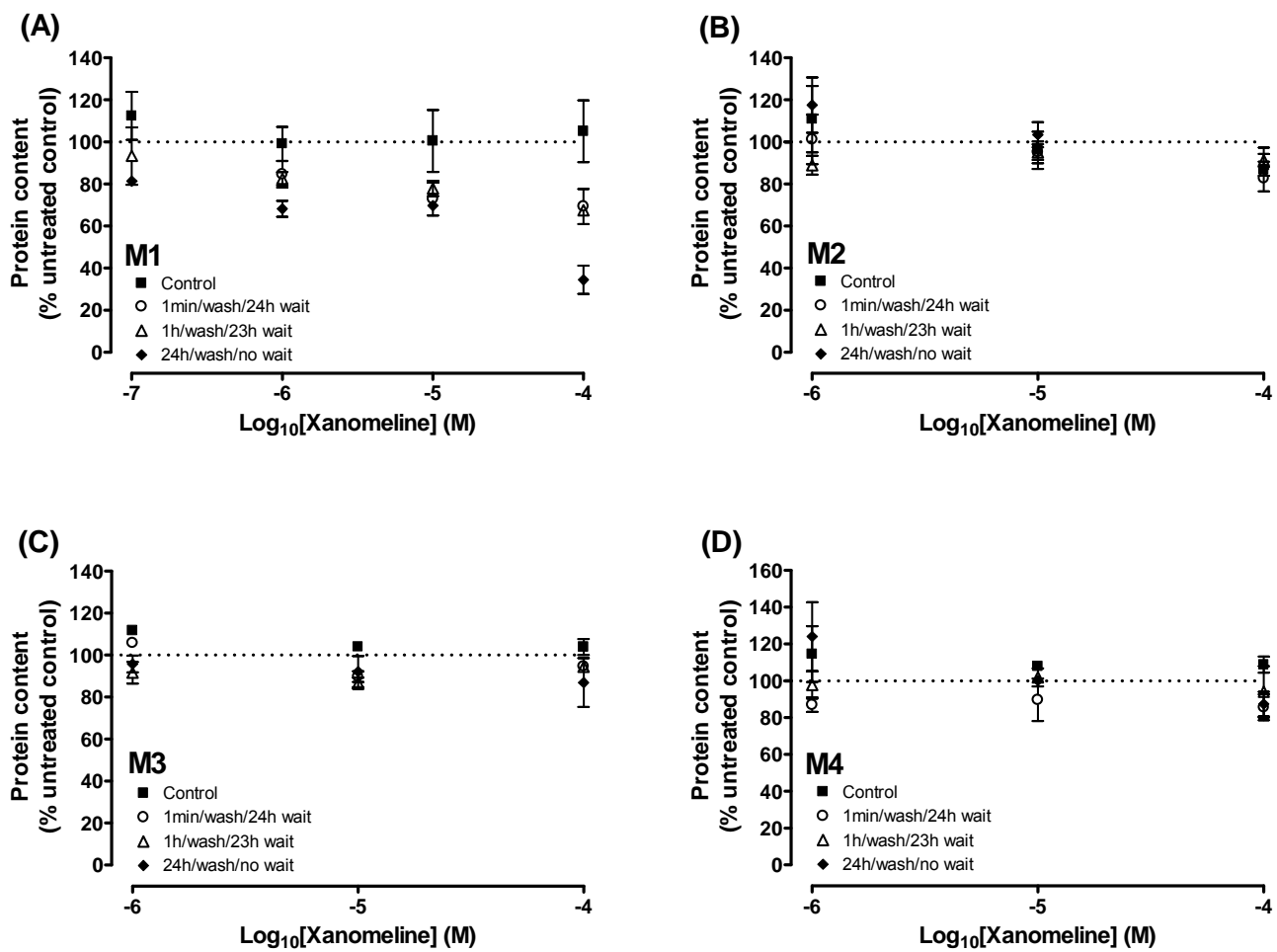
**Figure 11. Inhibition of [<sup>3</sup>H]NMS binding for long-term xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of xanomeline for 1 minute or 1 hour followed by washing and waiting 24 (open circles) or 23 (open triangles) hours, respectively. An additional group of cells was treated with xanomeline for 24 hours prior to washing (closed diamonds). Subsequently, cells were incubated with 0.2 nM [<sup>3</sup>H]NMS concurrently with xanomeline (closed squares) or following xanomeline pretreatments for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 13200, 12100, 21300 and 19000 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 3-9 experiments conducted in triplicate.

Figure 12



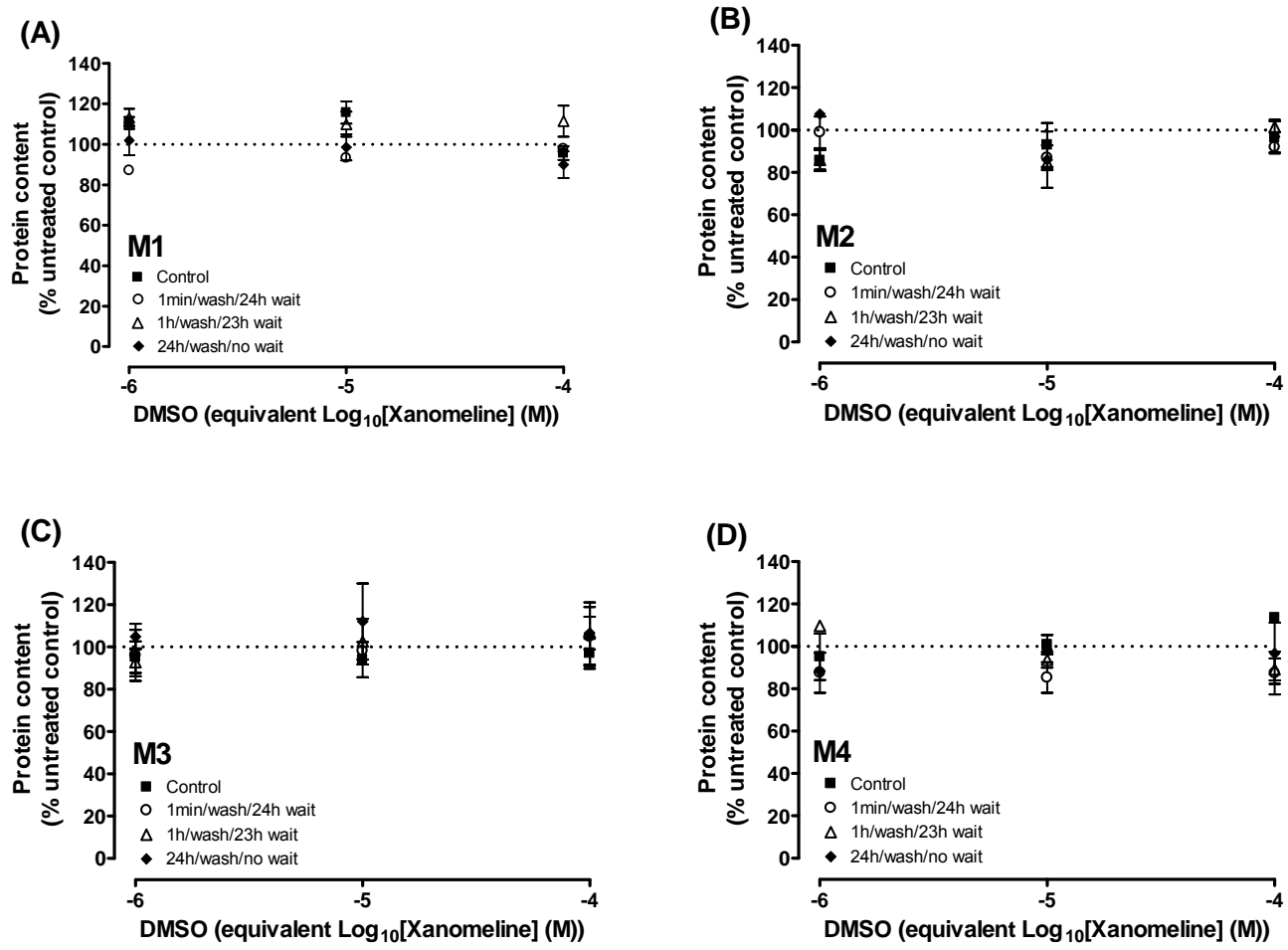
**Figure 12. Effects of long-term DMSO pretreatments on the inhibition of [<sup>3</sup>H]NMS binding in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of DMSO in media (corresponding to the concentrations used in the preparation of xanomeline) for 1 minute or 1 hour followed by washing and incubation in control media for 24 (open circles) or 23 (open triangles) hours, respectively. Additional cells were treated with DMSO for 24 hours prior to washing (closed diamonds). Subsequently, cells were exposed concurrently to DMSO and 0.2 nM [<sup>3</sup>H]NMS (closed squares) or to [<sup>3</sup>H]NMS following DMSO pretreatments for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control binding in the absence of DMSO treatment which had a mean of 11900, 15800, 21300 and 15400 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 2 experiments conducted in triplicate. Data from (A) provided by Marianne Grant.

Figure 13



**Figure 13. Effects of long-term xanomeline pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of xanomeline for 1 minute or 1 hour followed by washing and incubation in control media for 24 (open circles) or 23 (open triangles) hours, respectively. Additional cells were treated with xanomeline for 24 hours prior to washing (closed diamonds). Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with xanomeline in HEPES buffer for 1 hour (closed squares) or xanomeline pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of xanomeline treatment which had a mean of 130, 256, 153 and 196  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2-3 experiments conducted in triplicate.

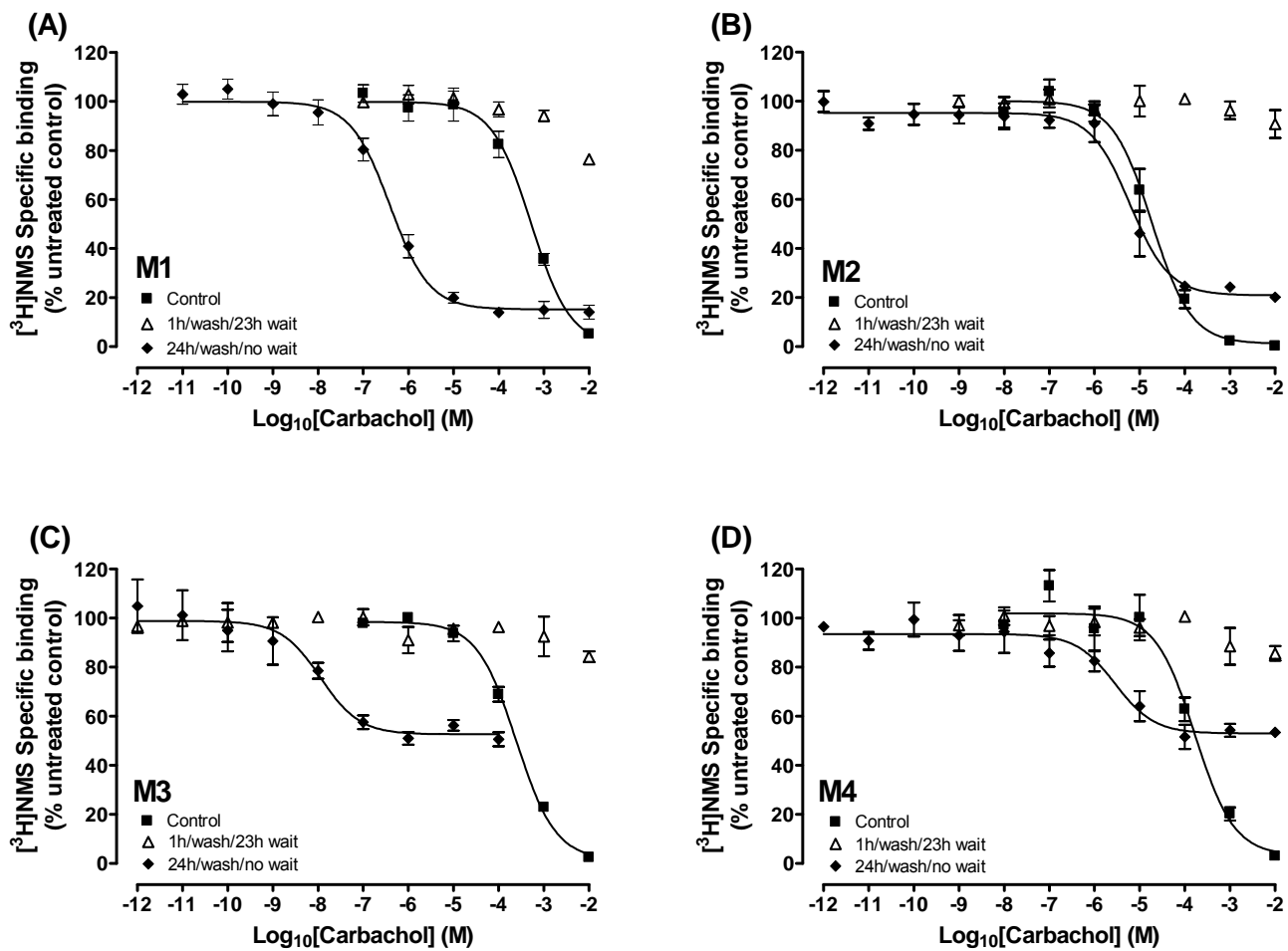
Figure 14



**Figure 14. Effects of long-term DMSO pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of DMSO in media (corresponding to the concentrations used in the preparation of xanomeline) for 1 minute or 1 hour followed by washing and incubation for 24 (open circles) or 23 (open triangles) hours, respectively. Additional cells were treated with DMSO for 24 hours prior to washing (closed diamonds). Subsequently, reactions were terminated using 1 M NaOH and protein content on untreated cells incubated with DMSO in HEPES buffer (closed squares) or DMSO pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of DMSO treatment which had a mean of 130, 256, 180 and 196  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

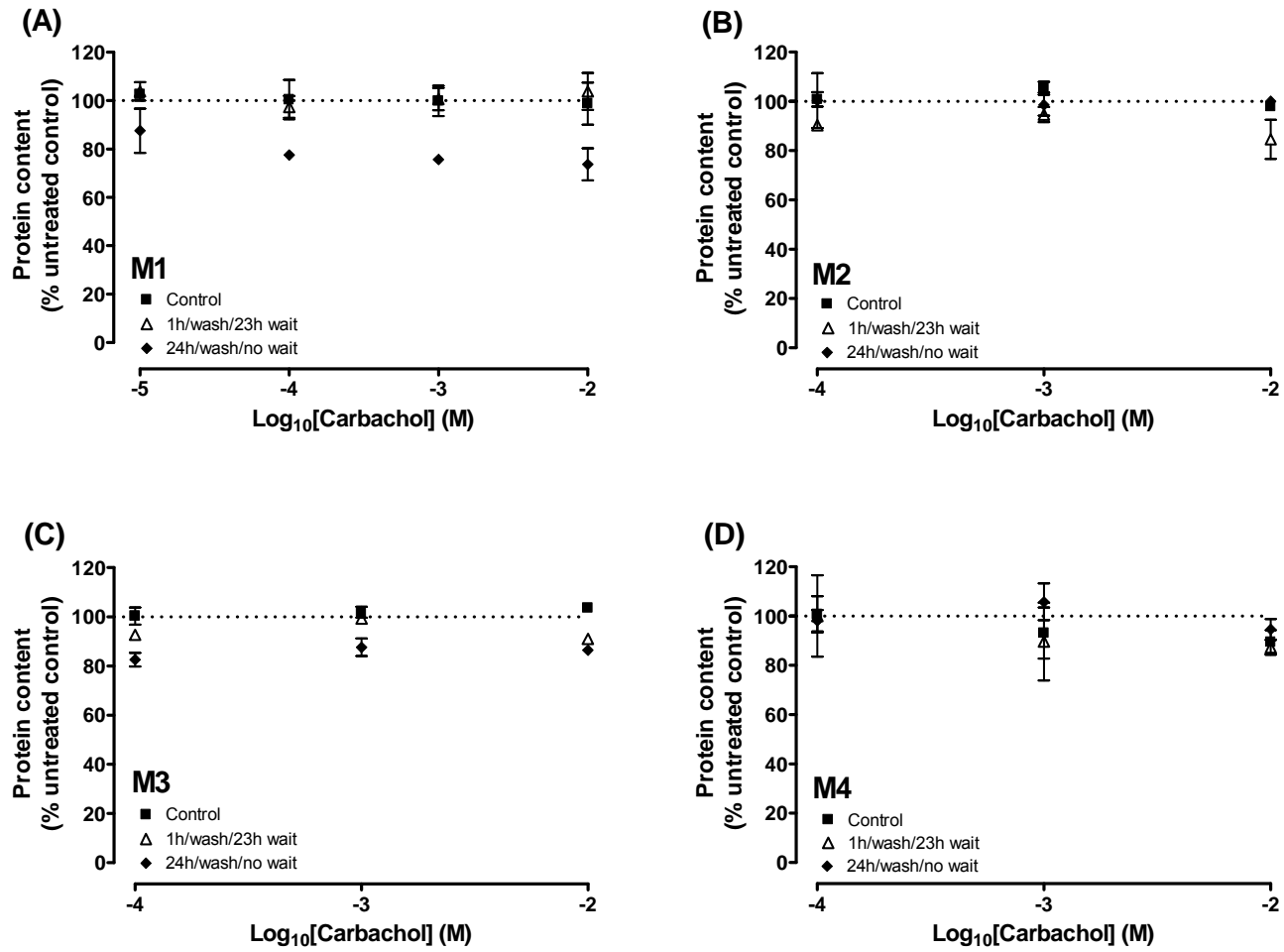


Figure 15



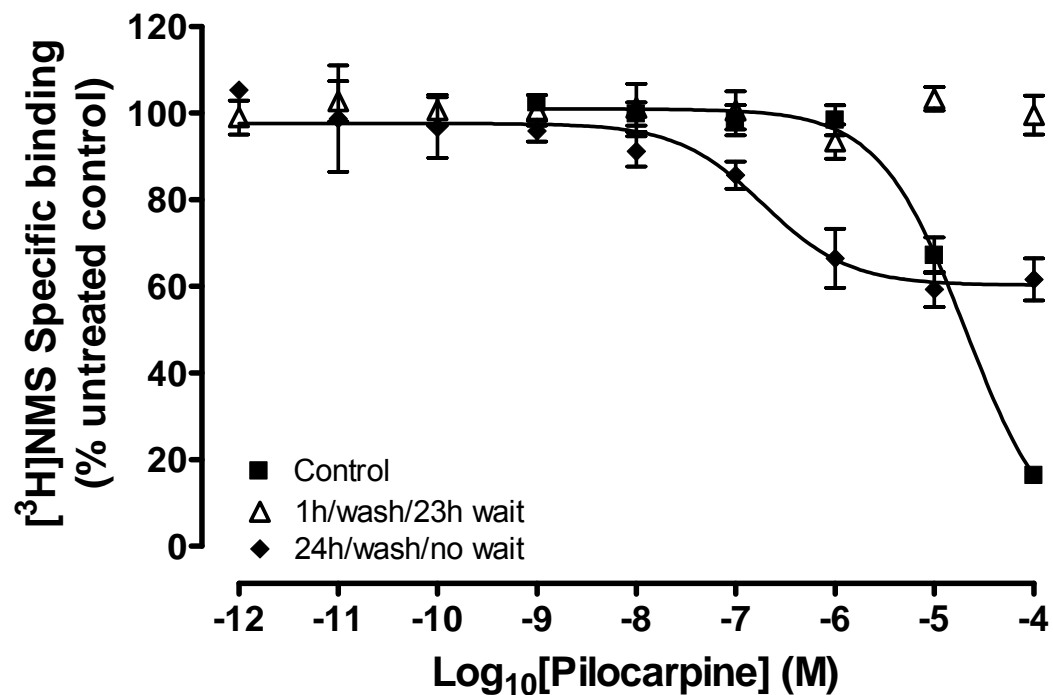
**Figure 15. Inhibition of [<sup>3</sup>H]NMS binding for long-term carbachol pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of carbachol for 1 hour followed by washing waiting 23 hours (open triangles). Additional cells were treated with carbachol for 24 hours prior to washing (closed diamonds). Subsequently, cells were incubated with 0.2 nM [<sup>3</sup>H]NMS concurrently with carbachol (closed squares) or following carbachol pretreatments for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Individual experiments were normalized to control binding in the absence of carbachol treatment which had a mean of 13000, 16500, 17700, and 16100 dpm/well for M1, M2, M3 and M4 receptors, respectively. Values represent the means ± S.E. of 3-6 experiments conducted in triplicate.

Figure 16



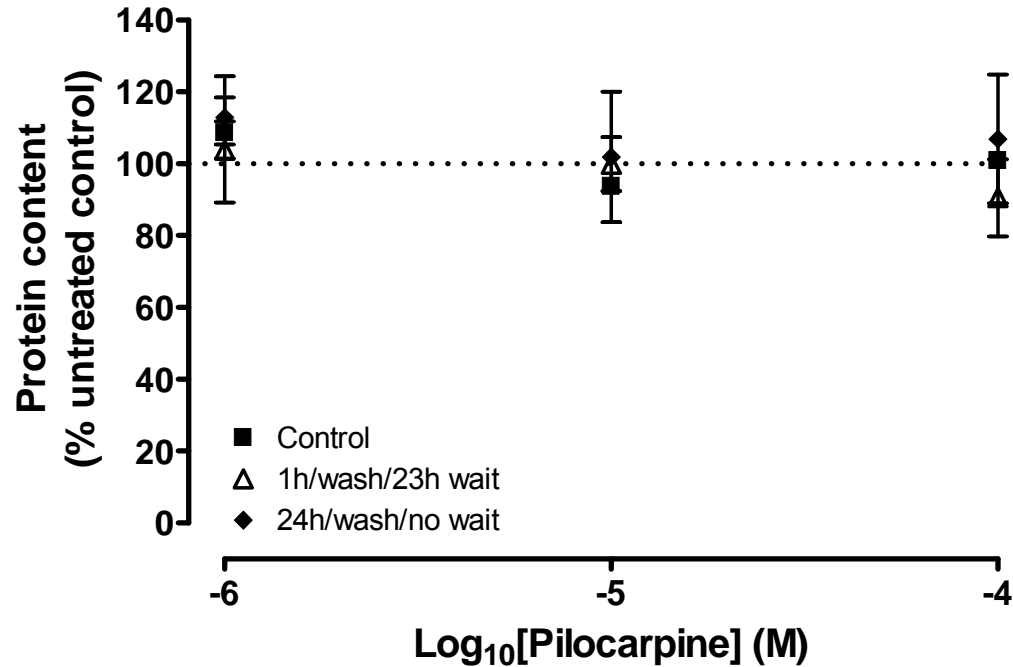
**Figure 16. Effects of long-term carbachol pretreatments on protein content in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with increasing concentrations of carbachol for 1 hour followed by washing and incubation in control media for 23 hours (open triangles). Additional cells were treated with carbachol for 24 hours prior to washing (closed diamonds). Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with carbachol in HEPES buffer for 1 hour (closed squares) or carbachol pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of carbachol treatment which had a mean of 180, 270, 153 and 296  $\mu\text{g}/\text{well}$  for M1, M2, M3 and M4 receptors, respectively. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

Figure 17



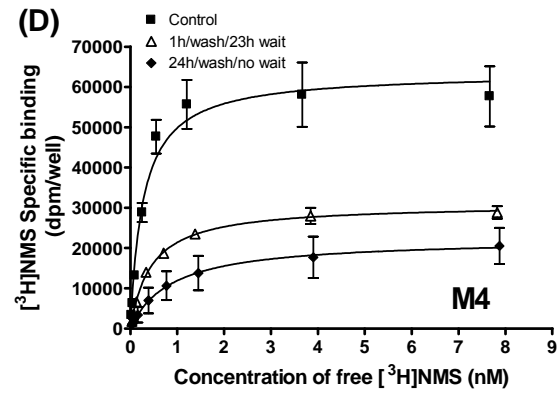
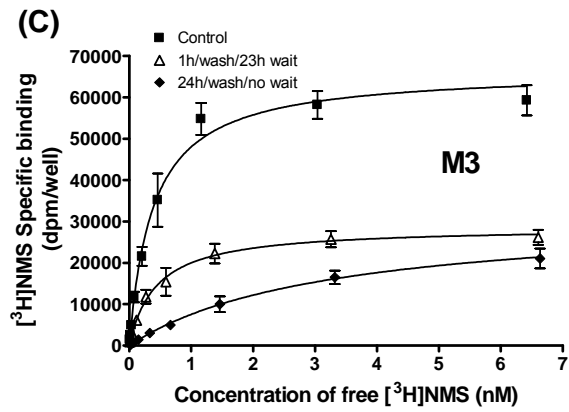
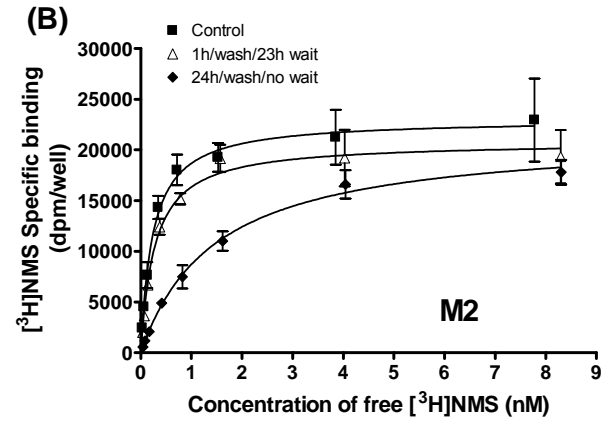
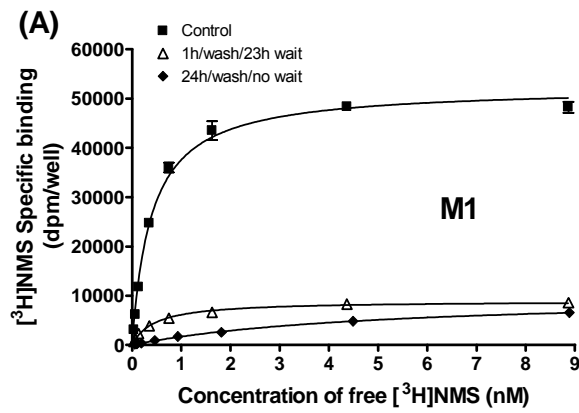
**Figure 17. Inhibition of [<sup>3</sup>H]NMS binding for long-term pilocarpine pretreatments in CHO cells stably expressing the hM3 receptor subtype.** Cells were pretreated with increasing concentrations of pilocarpine for 1 hour followed by washing waiting 23 hours (open triangles). Additional cells were treated with pilocarpine for 24 hours prior to washing (closed diamonds). Subsequently, cells were incubated with 0.2 nM [<sup>3</sup>H]NMS concurrently with pilocarpine (closed squares) or following pilocarpine pretreatments for 1 hour at 37°C. Individual experiments were normalized to control binding in the absence of pilocarpine treatment which had a mean of 17500 dpm/well. Values represent the means ± S.E. of 3 experiments conducted in triplicate.

Figure 18



**Figure 18. Effects of long-term pilocarpine pretreatments on protein content in CHO cells stably expressing hM3 receptor subtype.** Cells were pretreated with increasing concentrations of pilocarpine for 1 hour followed by washing and incubation in control media for 23 hours (open triangles). Additional cells were treated with pilocarpine for 24 hours prior to washing (closed diamonds). Subsequently, reactions were terminated using 1 M NaOH. Protein content on untreated cells incubated with pilocarpine in HEPES buffer for 1 hour (closed squares) or pilocarpine pretreated cells was determined using the method of Bradford (Bradford, 1976). A) M1 receptor, B) M2 receptor, C) M3 receptor and D) M4 receptor. Individual experiments were normalized to control protein in the absence of pilocarpine treatment which had a mean of 180  $\mu$ g/well. Values represent the means  $\pm$  S.E. of 2 experiments conducted in triplicate.

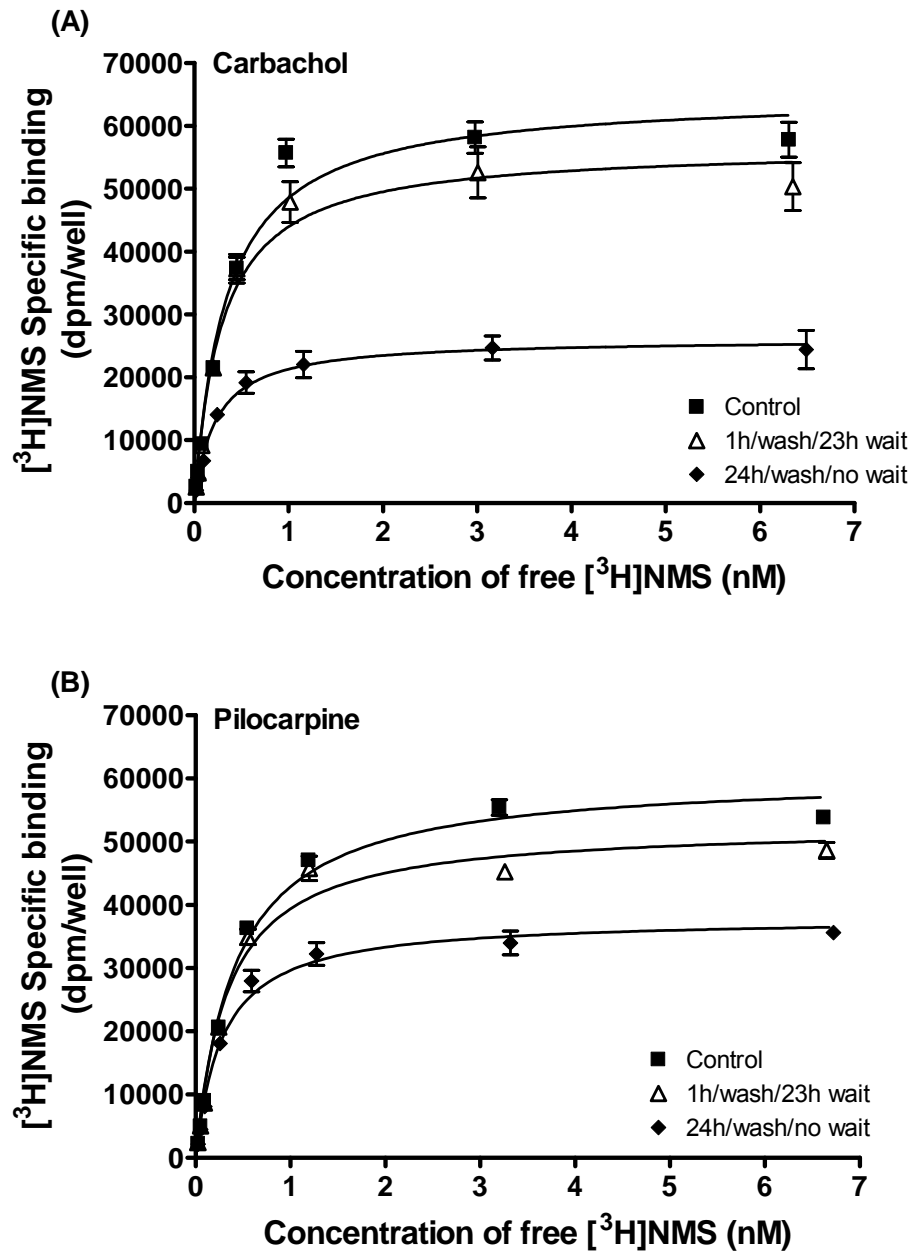
Figure 19



**Figure 19. [<sup>3</sup>H]NMS saturation binding subsequent to long-term xanomeline pretreatments in CHO cells stably expressing hM1-hM4 receptor subtypes.** Cells were pretreated with 10 μM xanomeline for 1 hour followed by washing and waiting 23 hours (open triangles). Additional cells were treated with xanomeline for 24 hours prior to washing (closed diamonds). Subsequently, untreated (closed squares) or xanomeline pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 8.9 nM) for 1 hour at 37°C. A) M1 receptor, B) M2 receptor, C) M3 receptor, and D) M4 receptor. Values represent the means ± S.E. of 3-4 experiments conducted in triplicate.



Figure 20



**Figure 20. [<sup>3</sup>H]NMS saturation binding subsequent to long-term agonist pretreatments in CHO cells stably expressing the hM3 receptor subtype.** Cells expressing the M3 receptor were exposed to 10  $\mu$ M agonist for 1 hour followed by washing and waiting 23 hours (open triangles). Additional cells were exposed to agonist for 24 hours prior to washing (closed diamonds). Subsequently, untreated (closed squares) or agonist pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.02 – 6.7 nM) for 1 hour at 37°C. A) Carbachol and B) Pilocarpine. Values represent the means  $\pm$  S.E. of 3 experiments conducted in triplicate.

**Table 1. Short- and long-term effects of xanomeline, carbachol or pilocarpine binding on [<sup>3</sup>H]NMS binding in CHO cells expressing hM1-hM4 receptor subtypes.** Cells were treated with increasing concentrations of xanomeline (1 pM – 100 μM), carbachol (1 pM – 10 mM) or pilocarpine (1 pM – 100 μM) for 1 minute or 1 hour followed by washing and immediate use in the binding assay or subsequent to incubation for 24 or 23 hours, respectively, in the absence of free agonist. Additional groups of cells were treated with increasing concentrations of agonists for 24 hours prior to washing. Pretreated and washed cells in all groups were incubated with 0.2nM [<sup>3</sup>H]NMS at 37°C for 1 hour. Control untreated cells were incubated simultaneously with agonist and 0.2nM [<sup>3</sup>H]NMS during the binding assay. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

	<b>M1</b>				<b>M2</b>	
	<b>pIC<sub>50</sub><sup>a</sup></b>	<b>pIC<sub>50</sub> – H<sup>b</sup></b>	<b>pIC<sub>50</sub> – L<sup>c</sup></b>	<b>I<sub>max</sub><sup>d</sup></b>	<b>pIC<sub>50</sub></b>	<b>I<sub>max</sub></b>
<b>Xanomeline</b>						
Control (untreated) <sup>e</sup>	6.5 ± 0.12			99 ± 0.3	6.9 ± 0.17	100 ± 0.3
1 min/washout/no wait	5.4 ± 0.13 <sup>§</sup>			85 ± 5.2 <sup>‡</sup>	5.0 ± 0.29 <sup>§</sup>	80 ± 6.0 <sup>‡</sup>
1 min/washout/24 h wait		8.2 ± 0.21 <sup>†</sup> (47 ± 4.3%)	4.6 ± 0.11 <sup>†</sup>	74 ± 6.2 <sup>‡</sup>	5.9 ± 0.44	41 ± 5.3 <sup>‡</sup>
1 h/washout/no wait	5.7 ± 0.05 <sup>§</sup>			97 ± 0.8	5.7 ± 0.12 <sup>§</sup>	87 ± 6.9
1 h/washout/23 h wait		8.6 ± 0.21* (43 ± 2.4%)	5.0 ± 0.06*	84 ± 2.8 <sup>‡</sup>	5.0 ± 0.15*	59 ± 3.0 <sup>‡</sup>
24 h/washout/no wait		9.8 ± 0.21* <sup>†</sup> (43 ± 3.7 %)	5.8 ± 0.06	97 ± 0.6	5.7 ± 0.12	93 ± 1.6
<b>Carbachol</b>						
Control (untreated)	3.3 ± 0.01			99 ± 0.3	4.7 ± 0.13	99 ± 0.8
1 h/washout/no wait	ND			ND	ND	ND
1 h/washout/23 h wait	ND			ND	ND	ND
24 h/washout/no wait	6.3 ± 0.08 <sup>#</sup>			88 ± 1.4 <sup>‡</sup>	5.2 ± 0.17	80 ± 1.4 <sup>‡</sup>

Table 1 cont.

	<b>M3</b>				<b>M4</b>			
	<b>pIC<sub>50</sub></b>	<b>pIC<sub>50</sub> - H</b>	<b>pIC<sub>50</sub> - L</b>	<b>I<sub>max</sub></b>	<b>pIC<sub>50</sub></b>	<b>pIC<sub>50</sub> - H</b>	<b>pIC<sub>50</sub> - L</b>	<b>I<sub>max</sub></b>
<b>Xanomeline</b>								
Control (untreated) <sup>e</sup>	7.1 ± 0.10			99 ± 0.4	7.1 ± 0.15			99 ± 0.7
1 min/washout/no wait	4.7 ± 0.08 <sup>s</sup>			93 ± 3.0	4.6 ± 0.18 <sup>s</sup>			85 ± 6.2 <sup>‡</sup>
1 min/washout/24 h wait		7.7 ± 0.75 <sup>†</sup> (40 ± 4.3%)	4.8 ± 0.43	82 ± 9.6 <sup>‡</sup>	4.8 ± 0.12			68 ± 3.1 <sup>‡</sup>
1 h/washout/no wait	5.6 ± 0.07 <sup>s</sup>			95 ± 0.9	5.9 ± 0.05 <sup>s</sup>			90 ± 2.6
1 h/washout/23 h wait		6.8 ± 0.39 (32 ± 9.9%)	4.3 ± 0.48	89 ± 5.5		7.1 ± 0.15* (38 ± 2.7%)	4.6 ± 0.10*	82 ± 1.1 <sup>‡</sup>
24 h/washout/no wait		9.0 ± 0.46* <sup>†</sup> (41 ± 3.4 %)	5.7 ± 0.06	94 ± 0.8		8.6 ± 0.29* <sup>†</sup> (29 ± 1.7 %)	5.9 ± 0.04 <sup>†</sup>	97 ± 0.6
<b>Carbachol</b>								
Control (untreated)	3.6 ± 0.05			98 ± 0.6	3.8 ± 0.05			97 ± 0.9
1 h/washout/no wait	ND			ND	4.7 ± 0.44			53 ± 4.5 <sup>‡</sup>
1 h/washout/23 h wait	ND			ND	ND			ND
24 h/washout/no wait	7.8 ± 0.19 <sup>#</sup>			48 ± 1.3 <sup>‡</sup>	5.5 ± 0.18 <sup>#</sup>			46 ± 2.7 <sup>‡</sup>
<b>Pilocarpine</b>								
Control (untreated)	4.7 ± 0.11			99 ± 0.9				
1 h/washout/no wait	ND			ND				
1 h/washout/23 h wait	ND			ND				
24 h/washout/no wait	6.7 ± 0.05 <sup>†</sup>			40 ± 5.4 <sup>‡</sup>				

Table 1 cont.

<sup>a</sup> Negative logarithm of the IC<sub>50</sub> for binding to a single potency site.

<sup>b</sup> Negative logarithm of the IC<sub>50</sub> for the high-potency site. Percent of binding sites shown in parentheses.

<sup>c</sup> Negative logarithm of the IC<sub>50</sub> for the low-potency site.

<sup>d</sup> Maximal percent inhibition of [<sup>3</sup>H]NMS binding.

<sup>e</sup> Control cells were incubated simultaneously with agonist and radioligand in the binding assay.

ND – parameters were not determined

<sup>§</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> between either the xanomeline 1min/wash/no wait or 1h/wash/no wait group compared to control.

<sup>†</sup> ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> where indicated between xanomeline 1min/wash/24h wait or 24h/wash/no wait pretreated groups compared to xanomeline 1min/wash/no wait treatment.

<sup>\*</sup> ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> where indicated between xanomeline 1h/wash/23h wait or 24h/wash/no wait pretreated groups compared to xanomeline 1h/wash/no wait treatment.

<sup>#</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> between carbachol 24h/wash/no wait compared to carbachol control.

<sup>±</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> between pilocarpine 24h/wash/no wait compared to pilocarpine control.

<sup>‡</sup> Significantly different ( $p < 0.5$ ) from 100 percent as determined by ANOVA followed by Dunnett's post-test comparison.

**Table 2. [<sup>3</sup>H]NMS saturation binding parameters for xanomeline, carbachol or pilocarpine pretreatment and waiting periods in CHO cells expressing hM1-hM4 receptor subtypes.** Cells were treated with 10 μM xanomeline, carbachol or pilocarpine for 1 hour followed by washing and immediate use in the binding assay or subsequent to incubation for 23 hours in the absence of free agonist. An additional group of cells was treated with agonist for 24 hours prior to washing. Untreated control or agonist pretreated cells were then incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 8.9 nM) for 1 hour at 37°C. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

	<b>M1</b>		<b>M2</b>		<b>M3</b>		<b>M4</b>	
	<b>K<sub>d</sub><sup>a</sup></b>	<b>B<sub>max</sub><sup>b</sup></b>	<b>K<sub>d</sub></b>	<b>B<sub>max</sub></b>	<b>K<sub>d</sub></b>	<b>B<sub>max</sub></b>	<b>K<sub>d</sub></b>	<b>B<sub>max</sub></b>
<b>Xanomeline</b>								
Control (untreated)	0.39 ± 0.02	52300 ± 700	0.24 ± 0.08	23200 ± 3200	0.42 ± 0.06	67700 ± 3200	0.27 ± 0.03	63600 ± 8400
1 h/washout/no wait	5.01 ± 0.18*	51000 ± 2500	1.42 ± 0.31*	26500 ± 5100	3.52 ± 0.52*	78500 ± 1600	2.89 ± 0.18*	64100 ± 5900
1 h/washout/23h wait	0.64 ± 0.10	9300 ± 300*	0.27 ± 0.05	20900 ± 2800	0.59 ± 0.21	30400 ± 2000*	0.43 ± 0.01	30400 ± 1200*
24 h/washout/no wait	5.11 ± 0.80*	10300 ± 1100*	1.47 ± 0.16*	21500 ± 1500	4.19 ± 0.93*	37000 ± 4600*	2.71 ± 0.26*	32900 ± 6500*
<b>Carbachol</b>								
Control (untreated)					0.34 ± 0.01	65000 ± 2900		
1 h/washout/23h wait					0.28 ± 0.01	56600 ± 4200		
24 h/washout/no wait					0.23 ± 0.03 <sup>†</sup>	26200 ± 2700 <sup>†</sup>		
<b>Pilocarpine</b>								
Control (untreated)					0.42 ± 0.03	60600 ± 960		
1 h/washout/23h wait					0.34 ± 0.02	52600 ± 850 <sup>§</sup>		
24 h/washout/no wait					0.28 ± 0.01 <sup>§</sup>	38000 ± 1500 <sup>§</sup>		

Table 2 cont.

<sup>a</sup> Equilibrium dissociation constant for [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell surface receptor density (dpm/well)

\*ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in  $K_d$  or  $B_{max}$  between the xanomeline pretreated groups compared with their untreated control for each of the individual subtypes.

† ANOVA followed by Dunnett's post-test detected a significant difference ( $p < 0.05$ ) in  $K_d$  and  $B_{max}$  between carbachol pretreated groups compared to carbachol control.

§ ANOVA followed by Dunnett's post-test detected a significant difference ( $p < 0.05$ ) in  $K_d$  and  $B_{max}$  between pilocarpine treated groups compared to pilocarpine control.

**CHAPTER 4: Determination of the acute and long-term functional effects of xanomeline at the M1 and M3 muscarinic receptors**

## Introduction

Muscarinic receptors are part of the G protein-coupled superfamily of receptors. Binding of an agonist to the receptor is thought to change the interaction of transmembrane domains 3, 5, 6 and 7 leading to the activation of down-stream signaling cascades (Eglen, 2005). This includes activation of phospholipase C, modulation of ion channels, activation of mitogen-activated protein kinase (involved in cell survival, differentiation and synaptic plasticity) and small GTPases (Volpicelli & Levey, 2004). Muscarinic M1 and M3 receptors are coupled to  $G_{q/11}$  G proteins through the third cytoplasmic domain. One way that the stimulation of the M1 or M3 receptors can be assessed is by determining the production of radiolabeled inositol phosphates following loading of the cells with radiolabeled *myo*-inositol (Berridge et al., 1983). Agonists can be classified based on their ability to stimulate a functional response. Full agonists elicit a maximal response, whereas partial agonists result in a submaximal response. Moreover, a partial agonist needs near or complete receptor occupancy to stimulate its maximal response (Hu & El-Fakahany, 1990).

We and others have demonstrated that xanomeline is able to bind to M1-M4 muscarinic receptor subtypes (Bymaster et al., 1997; Christopoulos et al., 1998; Grant & El-Fakahany, 2005)(Chapter 3). However, previous literature has proposed that xanomeline has functional selectivity for the M1 and M4 receptors (Shannon et al., 1994). Xanomeline has been shown to be a full agonist at the M1 receptor utilizing various assays of receptor function (Bymaster et al., 1997; Christopoulos et al., 1998). In contrast, previous research regarding the efficacy of xanomeline at the M3 receptor is contradictory. Bymaster et al. (1997) found that xanomeline was a partial agonist at the M3 receptor; whereas Wood et al. (1999) determined that xanomeline was a full agonist



at this receptor in their model systems. The discrepancy could be due to differences in receptor expression or the expression systems studied.

It is important to compare functional selectivity of xanomeline in the same type of cells that express individual subtypes of muscarinic receptors. This avoids complications caused by factors not related to the type of muscarinic receptor expressed, e.g. type and level of G proteins. Previously, we have shown that there is near equal receptor expression between the M1 and M3 receptors expressed in CHO cells as determined by saturation binding (Chapter 3 Fig. 10). It is also critical to compare the ability of long-term treatment with xanomeline to desensitize various subtypes of muscarinic receptors. Differential rates or extents of receptor desensitization would contribute to agonist selectivity under chronic treatment conditions. Thus, experiments were conducted to test the hypothesis that xanomeline has functional selectivity between the M1 and M3 receptors. We further extended the study to test the hypothesis that xanomeline induces differential long-term effects on subsequent agonist stimulation of M1 and M3 receptors.

## **Results**

### **Effects of xanomeline pretreatment on M1 and M3 receptor-mediated stimulation of inositol phosphate production**

Initial experiments were designed to compare the functional response elicited by xanomeline to that of the M3 full agonist carbachol and the partial agonist pilocarpine (Wang & El-Fakahany, 1993) in untreated cells. The maximal response stimulated by xanomeline was slightly lower than pilocarpine and approximately half that of carbachol, demonstrating that xanomeline displays properties of a partial agonist at the M3 receptor under our specific experimental conditions (Fig. 1 a). However, xanomeline exhibited a significantly ( $p < 0.05$ ) higher potency than either carbachol or pilocarpine (Table 1).

Wash-resistant xanomeline binding resulted in persistent activation of the M3 receptor (Fig. 1 b). Exposure of cells to xanomeline for one minute followed by washing led to a concentration-dependent increase in the hydrolysis of inositol phosphates, with a maximal response similar to that observed when it was continually present with untreated cells, although with a lower potency (Table 1). Interestingly, increasing the preincubation time with xanomeline to one hour prior to washing reduced the maximal response without a change in potency (Table 1).

The functional response of the M1 full agonist carbachol (Wang & El-Fakahany, 1993) was compared to that of xanomeline at the M1 receptor. In contrast to the M3 receptor, xanomeline had a similar maximal response to carbachol, supporting that it is a full agonist at the M1 receptor (Fig. 2 a). However, the potency of xanomeline was significantly ( $p < 0.05$ ) greater compared to carbachol. Results are summarized in Table 1.

Wash-resistant xanomeline was also able to simulate a functional response at the M1 receptor (Fig. 2 b). When cells were exposed to xanomeline for one minute followed by extensive washing there was a slight, but not significant, decrease in the maximal response compared to xanomeline treatment in the functional assay. In contrast, the same treatment resulted in a significant ( $p < 0.05$ ) decrease in potency. When xanomeline pretreatment time was increased to one hour it had no effect on the maximal response, but the potency was only slightly lower than when xanomeline was continually present in the binding assay.

## **Effects of acute xanomeline exposure on the ability of various muscarinic agonists to stimulate the production of inositol phosphates at the M3 and M1 receptors**

Brief pretreatment of cells expressing the M3 receptor with 10  $\mu$ M xanomeline followed by washing resulted in significant decreases in carbachol potency ( $p < 0.05$ ), with a more marked effect in the case of one hour pretreatment (Fig. 3 a; Table 2). No changes were observed in the maximal response elicited by carbachol following one minute or one hour pretreatment with xanomeline and washing (Fig. 3 a). The concurrent presence of xanomeline and carbachol during the functional assay resulted in a larger reduction in carbachol potency compared to that obtained following either pretreatment condition (Fig. 3 a; Table 2).

Further experiments were conducted to determine the effects of brief xanomeline treatment and washing on the response to partial agonists at the M3 receptor. Pretreatment with 10  $\mu$ M xanomeline for one minute or one hour followed by washing led to a significant ( $p < 0.05$ ) decrease in pilocarpine potency (by approximately 1 order of magnitude) (Fig. 3 b). This was accompanied by a significant ( $p < 0.05$ ) decrease in pilocarpine maximal response in the one hour-pretreatment group, but not the one-minute group (Fig. 3 b; Table 2). Xanomeline wash-resistant binding also resulted in antagonism of the response to a second addition of xanomeline. This was evidenced by a decrease in xanomeline potency without an alteration in efficacy (Fig. 3 c; Table 2).

Similar experiments were conducted at the M1 receptor to determine the ability of agonists to stimulate a functional response following acute xanomeline pretreatments. We noted that there were slight differences in the binding profiles when assays were performed on cells in suspension versus monolayer (discussed further in Appendix A). Thus, the functional response to carbachol stimulation was determined both in

monolayer and suspension following acute xanomeline pretreatments of cells in monolayer. When cells were treated with 300 nM xanomeline for one minute followed by washing there was no change in carbachol potency for cells assayed in monolayer (Fig. 4 a); however, there was a significant ( $p < 0.05$ ) increase in potency for the same pretreatment condition assayed in suspension (Fig. 4 b). Interestingly, there was a large increase in the basal production of inositol phosphates for cells assayed in suspension, but not for those assayed in monolayer. Neither assay condition resulted in a change in the maximal response to carbachol. When pretreatment time was increased to one hour there was also no change in carbachol maximal response (Fig. 4 c, d). However, there was a large increase in the basal production of inositol phosphates for both assay conditions. This large increase in basal response prevented the potency from being determined for cells assayed in suspension. Taken together, these results demonstrate that the response to carbachol stimulation following acute xanomeline pretreatments is similar, but not identical, for cells assayed in suspension compared to monolayer. Results are summarized in Table 3.

Further experiments were conducted in suspension at the M1 receptor to determine the ability of pilocarpine or xanomeline to stimulate a functional response following acute xanomeline pretreatments. Pretreatment with 300 nM xanomeline for one hour followed by washing did not affect that maximal response to either pilocarpine or xanomeline (Fig. 4 c, d). In contrast, one hour pretreatment with xanomeline resulted in a significant ( $p < 0.05$ ) decrease in xanomeline potency, but not pilocarpine potency compared to their respective controls. Results are summarized in Table 3.

### **Effects of acute agonist exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the M3 receptor**

To determine if the acute effects of xanomeline on agonist stimulation were unique to xanomeline, cells expressing the M3 receptor were exposed to 10  $\mu$ M pilocarpine for one hour followed by washing and stimulation with a second application of pilocarpine. Under these conditions, there was no change in the potency of pilocarpine and only a slight decrease in the maximal response compared to pilocarpine stimulation in untreated cells (Fig. 5). Results are summarized in Table 2.

### **Stimulation of inositol phosphates production using exogenous phospholipase C subsequent to short- or long-term treatment with xanomeline at the M3 and M1 receptors**

In order to demonstrate that the pool of labeled polyphosphoinositides at either the M3 or M1 receptor was not depleted following short- or long-term xanomeline pretreatments, cells were stimulated with exogenous phospholipase C following the various pretreatments with xanomeline. No differences in the production of inositol phosphates were observed in any of the experimental groups for short-term xanomeline treatments at the M1 (Fig. 6 a) or M3 (Fig. 6 b) receptors or for the long-term xanomeline treatments at the M1 (Fig. 7 a) or M3 (Fig. 7 b) receptors.

### **Effects of long-term xanomeline exposure on the ability of various muscarinic agonists to stimulate inositol phosphate production at the M3 and M1 receptors**

The full agonist carbachol was utilized to determine the long-term effects of xanomeline pretreatment on the sensitivity of the M3 receptor. Pretreatment with 10  $\mu$ M xanomeline for one minute or one hour followed by washing and waiting for a day

resulted in nearly identical concentration response curves, displaying significant ( $p < 0.05$ ) decreases in the potency of carbachol without significant changes in maximal response compared to control (Fig. 8 a; Table 2). A more marked decrease in carbachol potency (2.4 orders of magnitude) was observed when cells were pretreated with 10  $\mu\text{M}$  xanomeline for 24 hours before washing (Fig. 8 a; Table 2). Again, there was no change in carbachol maximal response.

Lack of demonstrable effects of xanomeline pretreatment on the maximal response elicited by the full agonist carbachol, in spite of the observed large decrease in the number of available cell-surface receptors (Chapter 3 Fig. 19), might be due to the involvement of spare receptors. We therefore tested the effects of xanomeline on the receptor response to partial agonists, as such ligands require full receptor occupancy to elicit maximal receptor activation. Cells were pretreated with 10  $\mu\text{M}$  xanomeline for 24 hours prior to washing, or preincubated with 10  $\mu\text{M}$  xanomeline for either one minute or one hour followed by washing and prolonged waiting in ligand-free media. In all cases, comparable significant ( $p < 0.05$ ) decreases in the maximal response to the partial agonist pilocarpine were observed (Fig. 8 b). These effects were accompanied by a significant ( $p < 0.05$ ) reduction in pilocarpine potency, with a more pronounced effect in case of the 24-hour pretreatment group (Fig. 8 b; Table 2). Of particular note, pretreatment with xanomeline resulted in more marked attenuation of its own stimulatory effects as compared to those of carbachol and pilocarpine (Fig. 8 c). However, xanomeline maximal receptor activation and potency under these conditions could not be quantified since a full concentration response curve could not be attained, due to the concentration limitations presented by using DMSO as a solvent.

As mentioned previously and discussed further in Appendix A, we observed a difference between the results of binding assays conducted in monolayer versus

suspension, so functional assays assessing the ability of 300 nM carbachol to stimulate the production of inositol phosphates subsequent to long-term xanomeline pretreatments were performed both in monolayer and in suspension at the M1 receptor. When cells expressing the M1 receptor were treated with 300 nM xanomeline for one minute or one hour followed by washing and overnight incubation in ligand free media there was no change in the ability of carbachol to stimulate a maximal response for cells assayed in monolayer (Fig. 9 a) or suspension (Fig. 9 b). However, there was a significant ( $p < 0.05$ ) decrease in potency in suspension and a trend in monolayer towards a decrease in the potency following one hour, but not one minute pretreatment in either case. The response to carbachol stimulation following treatment with xanomeline for 24 hours prior to washing was also similar for both monolayer (Fig. 9 a) and suspension (Fig. 9 b) assay groups with a decrease in both potency and maximal stimulation. These results demonstrate that the ability of carbachol to stimulate the production of inositol phosphates following long-term pretreatment with xanomeline is similar between monolayer and suspension assays. Results are summarized in Table 3.

Further experiments were conducted with pilocarpine and xanomeline stimulation in suspension subsequent to long-term xanomeline pretreatments in cells expressing the M1 receptor. In our expression system, pilocarpine and xanomeline are both full agonists at the M1 receptor. When cells were pretreated with 300 nM xanomeline for one hour followed by washing and overnight incubation or treated for 24 hours prior to washing there was a significant ( $p < 0.05$ ) decrease in the maximal stimulation of inositol phosphates in response to pilocarpine, along with a significant ( $p < 0.05$ ) decrease in pilocarpine potency (Fig. 9 c), with a more pronounced effect following 24 hour pretreatment. Similar results were obtained with xanomeline stimulation following 24 hour xanomeline pretreatments. However, xanomeline pretreatment for one hour

followed by washing and waiting 23 hours resulted in only a moderate decrease in the potency of xanomeline and a smaller, yet significant ( $p < 0.05$ ), decrease in maximal stimulation (Fig. 9 d) compared to xanomeline stimulation alone. Results are summarized in Table 3.

### **Effects of long-term carbachol or pilocarpine exposure on the ability of agonists to stimulate inositol phosphate production at the M3 and M1 receptors**

Experiments were conducted to ascertain if the observed delayed effects of wash-resistant xanomeline binding on receptor function at the M3 and M1 receptors (Figs. 8 & 9) were unique to xanomeline or due to general effects elicited by long-term exposure to any muscarinic agonist. A 10  $\mu$ M concentration of either carbachol or pilocarpine was selected since at this concentration both agonists produce maximal receptor activation and maximal inhibition of [ $^3$ H]NMS binding following prolonged exposure.

When cells expressing the M3 receptor were pretreated with carbachol or pilocarpine for one hour followed by washing and incubation in control media for 23 hours, no change in efficacy or potency of carbachol in stimulating phosphoinositide hydrolysis was observed (Figs. 10 b, c; Table 2). In contrast, pretreatment with carbachol for 24 hours followed by washing resulted in a significant ( $p < 0.05$ ) decrease in the maximal response and potency of a second addition of carbachol (Fig. 10 b; Table 2). A similar long-term treatment with pilocarpine did not alter either concentration-response parameters of carbachol (Fig. 10 c; Table 2). In addition, pretreatment with pilocarpine for one hour followed by washing and waiting did not modify the concentration-response relationship of pilocarpine. However, pilocarpine treatment for 24 hours prior to washing followed by a second addition of pilocarpine demonstrated



significant ( $p < 0.05$ ) decreased potency as well as a marked decrease in efficacy (Fig. 10 d), similar to what was observed with carbachol pretreatment followed by carbachol stimulation (Fig. 10 b).

Similar experiments utilizing carbachol stimulation following carbachol pretreatment were conducted in cells expressing the M1 receptor. When cells were pretreated with 10  $\mu$ M carbachol for one hour followed by overnight incubation in ligand free media there was no difference in the potency or efficacy following carbachol stimulation compared to control (Fig. 10 a). When cells were pretreated with carbachol for 24 hours prior to washing there was no change in efficacy following a second addition of carbachol to stimulate the production of inositol phosphates compared to carbachol stimulation alone (Fig. 10 a). In contrast, the same treatment resulted in a substantial decrease in the potency of carbachol (Fig. 10 a). Results are summarized in Table 3.

Further experiments were performed to determine the effects of 24 hour carbachol pretreatment on the ability of xanomeline to stimulate a response at either the M3 or M1 receptor (Fig. 11). Pretreatment of cells expressing the M3 receptor for 24 hours with carbachol followed by washing and stimulation with increasing concentrations of xanomeline resulted in minimal production of inositol phosphates observed only at the highest concentration of xanomeline used (100  $\mu$ M) (Fig. 11 b; Table 2). Pretreatment of cells expressing the M1 receptor for 24 hours with carbachol prior to xanomeline stimulation also resulted in minimal production of inositol phosphates (Fig. 11 a; Table 3). However, the response occurred with lower concentrations of xanomeline than needed at the M3 receptor.

## **Discussion**

The functional consequences of xanomeline treatment were studied by determining the accumulation of inositol phosphates (Figs. 1 & 2). Xanomeline was found to be a potent partial agonist at the M3 muscarinic receptor (Fig. 1) and a potent full agonist at the M1 muscarinic receptor (Fig. 2). This is in agreement with previous studies (Bymaster et al., 1997; Christopoulos et al., 1998). However, Wood et al. (1999) showed that xanomeline acted as a full agonist at M3 receptors in their expression system. This discrepancy may be due to different receptor expression levels or variation in the functional assays employed to assess receptor sensitivity. The wash-resistant component of xanomeline binding established after exposure to the drug for one minute was able to stimulate a concentration-dependent functional response at both the M3 and M1 receptor. The maximal response at the M3 receptor following one-minute xanomeline exposure and washing was similar to that observed when xanomeline was present during the functional assay, albeit with a lower potency. In contrast, there was a slight decrease in efficacy accompanied by a decrease in potency following one minute xanomeline treatment and washing compared to the presence of xanomeline during the functional assay at the M1 receptor. When pretreatment with xanomeline was increased to one hour followed by washing it resulted in a slightly lower efficacy compared to xanomeline continuously present in the assay at the M3 and M1 receptors. This time-dependent reduction in receptor sensitivity might be due to xanomeline-induced desensitization of the receptor. It has been demonstrated that an agonist can cause desensitization without changing receptor expression (Maloteaux & Hermans, 1994).

The notion that desensitization was occurring without changing cell-surface receptor density was tested further by examining the effects of xanomeline treatment on the ability of other agonists to stimulate the production of inositol phosphates (Figs. 4 &

5). When xanomeline was present for one minute or one hour followed by washing and immediate use it decreased the potency of carbachol without reduction of its maximal response at both the M3 and M1 receptors. These results are in agreement with saturation binding experiments in which there was marked reduction in the affinity of the radioligand for either receptor in the absence of change in receptor density. However, at the M3 receptor, pretreatment with xanomeline for one hour resulted in wash-resistant attenuation of not only the potency of the partial agonist pilocarpine, but also its maximal response. This might be due to the higher susceptibility of partial agonists to changes in receptor sensitivity (Hu et al., 1991). Alternatively, these results could be explained by differential allosteric modulation by xanomeline of the physical interaction of different agonists with the receptor (Lanzafame et al., 2006; May et al., 2005). In contrast, pretreatment with xanomeline for one hour followed by washing did not have an effect on potency or efficacy of pilocarpine stimulation at the M1 receptor. This may be because pilocarpine is a full agonist at the M1 receptor. At either the M3 or M1 receptor, pretreatment with xanomeline followed by subsequent stimulation with itself resulted in a reduction of the potency with no change in efficacy. Interestingly, xanomeline and pilocarpine are both partial agonists at the M3 receptor; however, the effects of xanomeline pretreatment prior to their stimulation are slightly different, further suggesting allosteric modulation of the receptor by xanomeline.

The long-term effects of xanomeline pretreatment are more marked than the acute effects. When cells expressing the M3 receptor were treated with xanomeline for one minute or one hour followed by washing and prolonged waiting there was a decrease in the potency of both carbachol and pilocarpine, albeit the shift in potency was less pronounced than that observed following acute xanomeline pretreatments (Fig. 8). This corresponds well to the observed reversal of the effects of xanomeline on [<sup>3</sup>H]NMS

affinity following prolonged incubation of pretreated and washed cells in control media. In contrast, there was a greater decrease in the potency of both carbachol and pilocarpine following brief xanomeline treatment prior to washing and waiting at the M1 receptor, compared to acute xanomeline pretreatments (Fig. 9). This discrepancy could be a result of differential modulation of the receptor between subtypes. In contrast, wash-resistant xanomeline binding followed by waiting at both the M3 and M1 receptors resulted in additional delayed reduction in the maximal functional response elicited by carbachol and pilocarpine that is likely due to reduction in cell-surface receptor density detected in radioligand binding experiments. Pretreatment with xanomeline for 24 hours prior to washing resulted in a more pronounced effect on the potency of carbachol and pilocarpine compared to brief xanomeline pretreatment followed by washing and waiting at both subtypes. This might be due to the larger decrease in receptor potency detected in radioligand binding studies under the latter treatment conditions. Something else might be happening at the M1 receptor since the decrease in radioligand affinity is not greater for 24 hours. In contrast, when cells expressing the M3 receptor were briefly treated with xanomeline followed by washing and further incubation or treated for 24 hours prior to washing it markedly reduced the ability of xanomeline to stimulate a subsequent response. This reduction was to a greater extent than was seen with other agonists, suggesting the possibility of differential allosteric modulation of the receptor by xanomeline. However, at the M1 receptor, the reduction in the ability of xanomeline to stimulate a response following pretreatment with itself was not as attenuated. This further suggests a slight difference in xanomeline regulation between the M1 and M3 receptors.

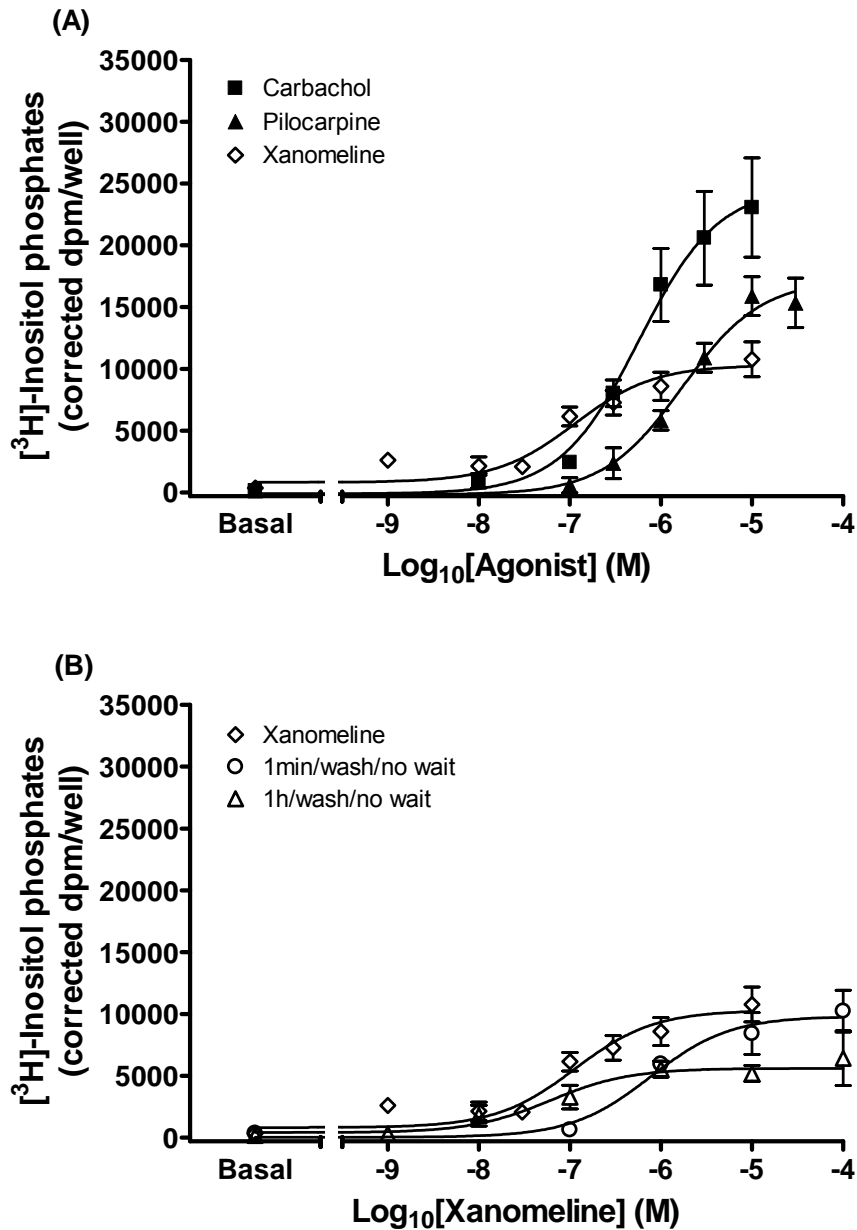
The observed delayed effects of brief xanomeline wash-resistant binding were associated with attenuation of the receptor response to activation by carbachol. Control

experiments with the conventional agonists carbachol and pilocarpine confirmed that the functional changes observed following xanomeline pretreatments were due to unique ligand-specific characteristics rather than being a general property of agonist binding to the receptor (Fig. 10). When cells expressing the M3 receptor were pretreated with pilocarpine for one hour followed by washing and waiting 23 hours or for 24 hours prior to washing, no changes in carbachol-induced receptor stimulation were observed. Similarly, pretreatment with carbachol for one hour followed by washing and waiting 23 hours did not alter the ability of carbachol to elicit a response at either the M3 or M1 receptors. However, when cells expressing either the M3 or M1 receptor were treated for 24 hours with carbachol prior to washing, there was a depression in the maximal response in addition to a reduction in potency of the carbachol-mediated response (Fig. 10). Literature supports that the observed decrease in the functional response may be due to internalization of cell-surface receptors in addition to uncoupling of the receptor from G proteins (Hu et al., 1991; Yang et al., 1993). Pretreatment of cells expressing the M3 receptor with pilocarpine produced changes in its ability to subsequently activate the receptor similar to those described above for carbachol. Contradictory reports of receptor regulation by pilocarpine exist. Previous research has shown that prolonged exposure to pilocarpine results in a decrease in cell-surface receptor density (Shifrin & Klein, 1980). In contrast, another report suggested that pilocarpine has little effect on the actual number of receptors, as the apparent down-regulation was due to pilocarpine being released when cells were lysed prior to the binding assay (Milligan & Strange, 1983). However, in our paradigm 24-hour exposure of cells expressing the M3 receptor to pilocarpine prior to washing resulted in a decrease in cell-surface receptor density (Fig. 10). Thus, the change in the functional response to pilocarpine can be attributed to a decrease in cell-surface receptor density following continuous agonist exposure,

because a partial agonist needs near or full receptor occupancy to stimulate a maximal response (Hu & El-Fakahany, 1990). Additionally, 24-hour pretreatment with carbachol resulted in a larger reduction in the ability of xanomeline to stimulate a functional response at either the M3 or M1 receptors compared to stimulation by other agonists. Thus, these results further demonstrate that xanomeline interacts in a unique manner with muscarinic receptors.

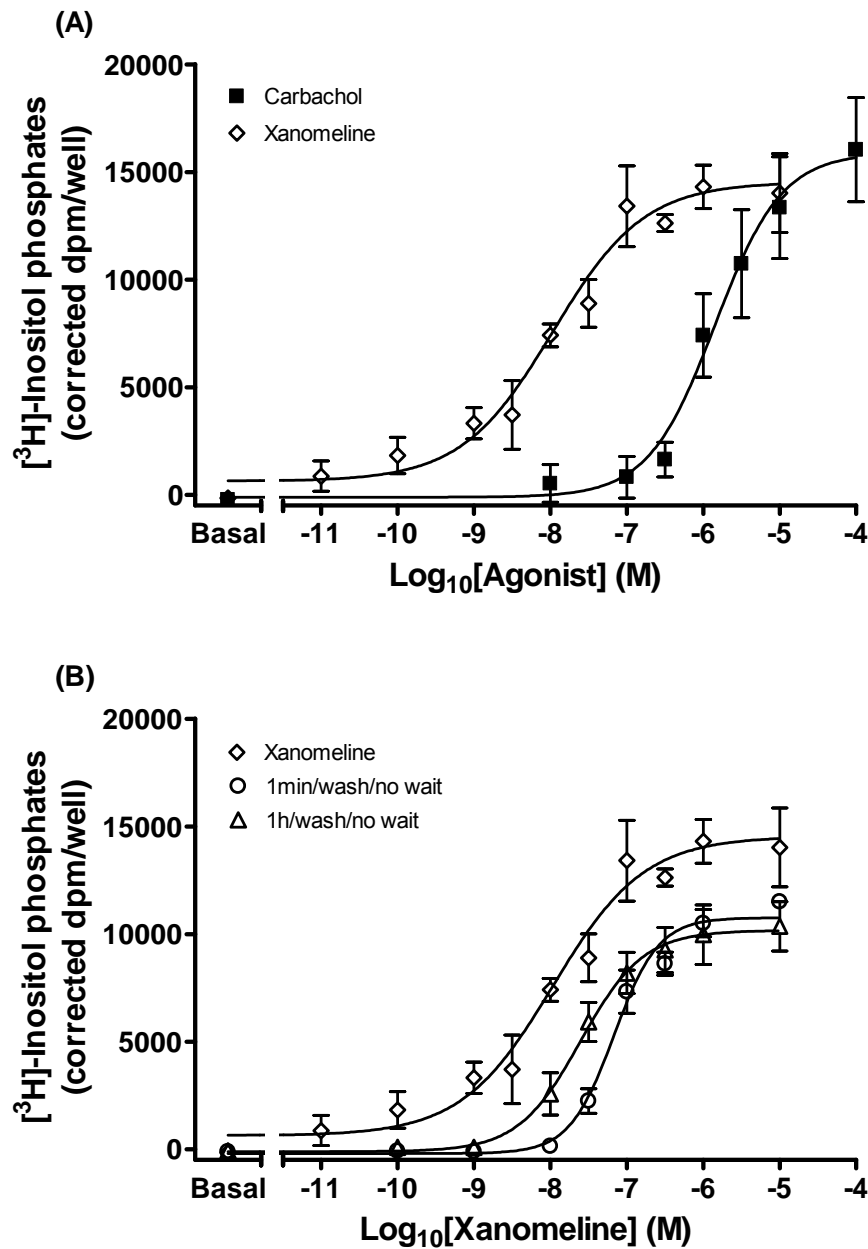
In summary, we have demonstrated that xanomeline is a potent partial agonist at the M3 receptor and a potent full agonist at the M1 receptor. Additionally, wash-resistant xanomeline can stimulate a functional response at both receptor subtypes. Acute pretreatment with xanomeline is able to modulate the response to subsequent stimulation by various agonists. Similarly, prolonged exposure to xanomeline results in antagonism of the response to subsequent stimulation with agonists. This antagonism was slightly more pronounced at the M3 receptor compared to the M1 receptor. These effects are unique to xanomeline rather than being a general property of conventional agonists. These results are in agreement with the slight differences observed between the M1 and M3 receptor in radioligand binding studies. Furthermore, these results demonstrate that xanomeline is able to modulate receptor function through binding at an allosteric site on the receptor.

Figure 1



**Figure 1. hM3 receptor activation by xanomeline reversible and wash-resistant binding.** A) Cells were incubated at 37°C for 1 hour with increasing concentrations of carbachol (closed squares), pilocarpine (closed triangles) or xanomeline (open diamonds) in the presence of 10 mM lithium chloride. B) Cells were pretreated with increasing concentrations of xanomeline for 1 minute (open circles) or 1 hour (open diamonds) followed by washing and determination of the accumulation of inositol phosphates by wash-resistant xanomeline. In the third group (open diamonds) xanomeline was only present during determination of inositol phosphates accumulation. Values represent the means  $\pm$  S.E. of 3-5 experiments conducted in triplicate.

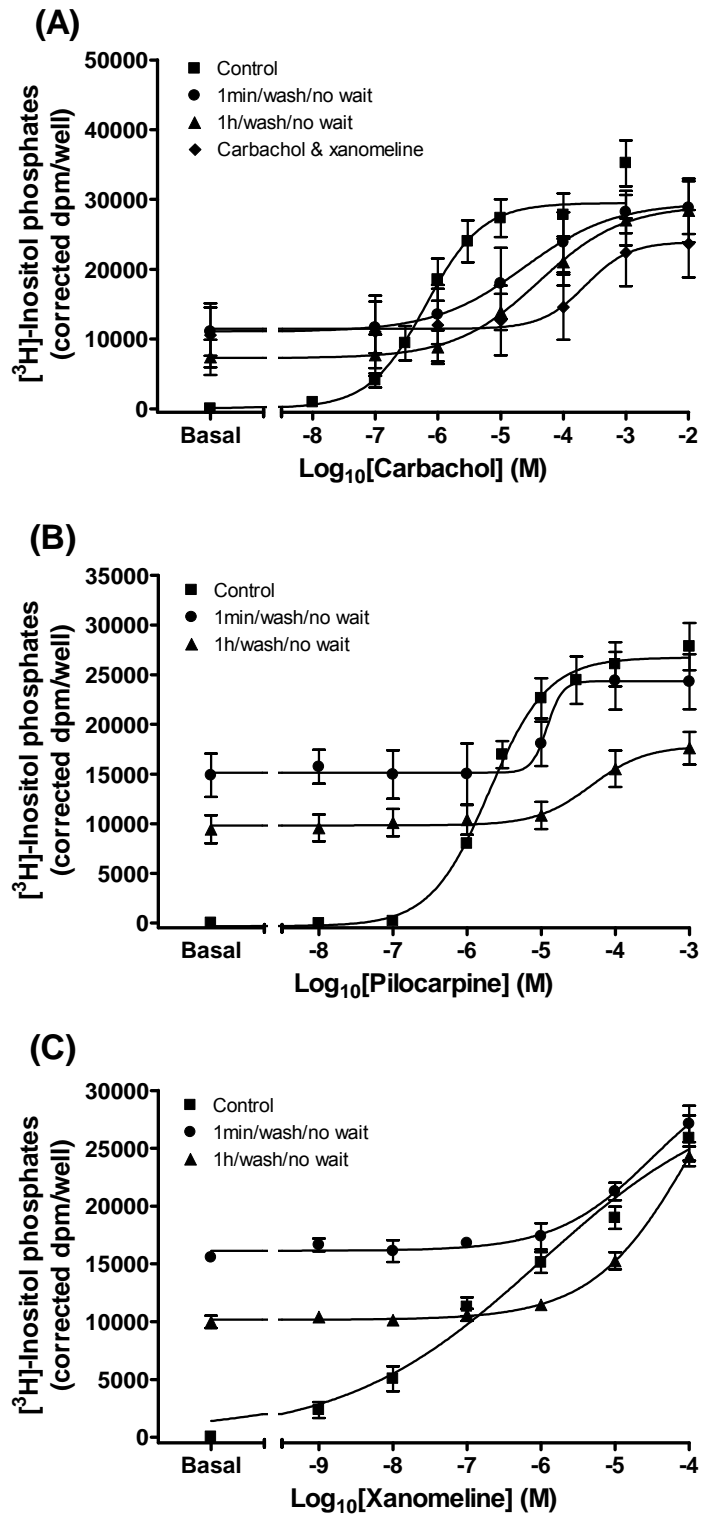
**Figure 2**



**Figure 2. hM1 receptor activation by xanomeline reversible and wash-resistant binding.** A) Cells were incubated at 37°C for 1 hour with increasing concentrations of carbachol (closed squares) or xanomeline (open diamonds) in the presence of 10 mM lithium chloride. B) Cells were pretreated with increasing concentrations of xanomeline for 1 minute (open circles) or 1 hour (open diamonds) followed by washing and determination of the accumulation of inositol phosphates by wash-resistant xanomeline. In the third group (open diamonds) xanomeline was only present during determination of inositol phosphates accumulation. Values represent the means  $\pm$  S.E. of 3-4 experiments conducted in triplicate. Data provided by Marianne Grant.

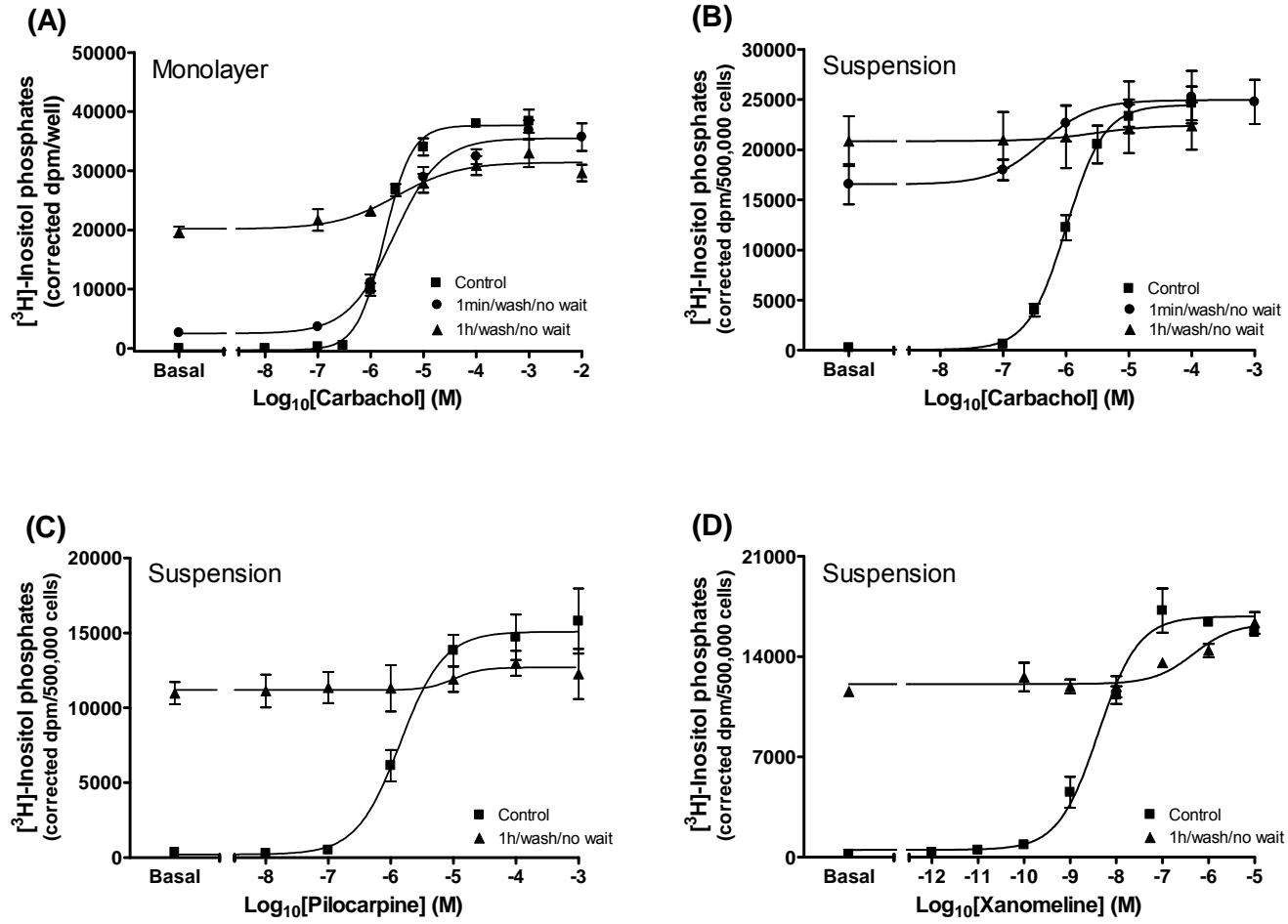


Figure 3



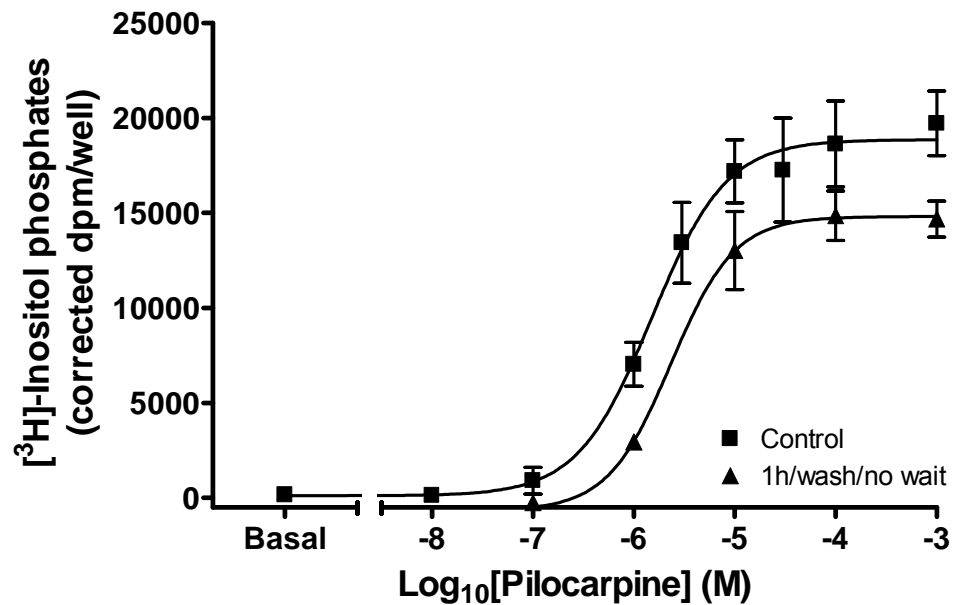
**Figure 3. Effects of acute xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM3 receptor.** Cells were pretreated with 10  $\mu$ M xanomeline for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing and immediate use. Subsequently, cells were treated with increasing concentrations of A) carbachol, B) pilocarpine, or C) xanomeline and the accumulation of inositol phosphates was determined over a 1-hour incubation period at 37°C in the presence of 10 mM lithium chloride for untreated cells (closed squares) and those pretreated with xanomeline. A) An additional group of cells was simulated simultaneously with increasing concentrations of carbachol and 10  $\mu$ M xanomeline (closed diamonds). Values represent the means  $\pm$  S.E. of 3-7 experiments conducted in triplicate.

Figure 4



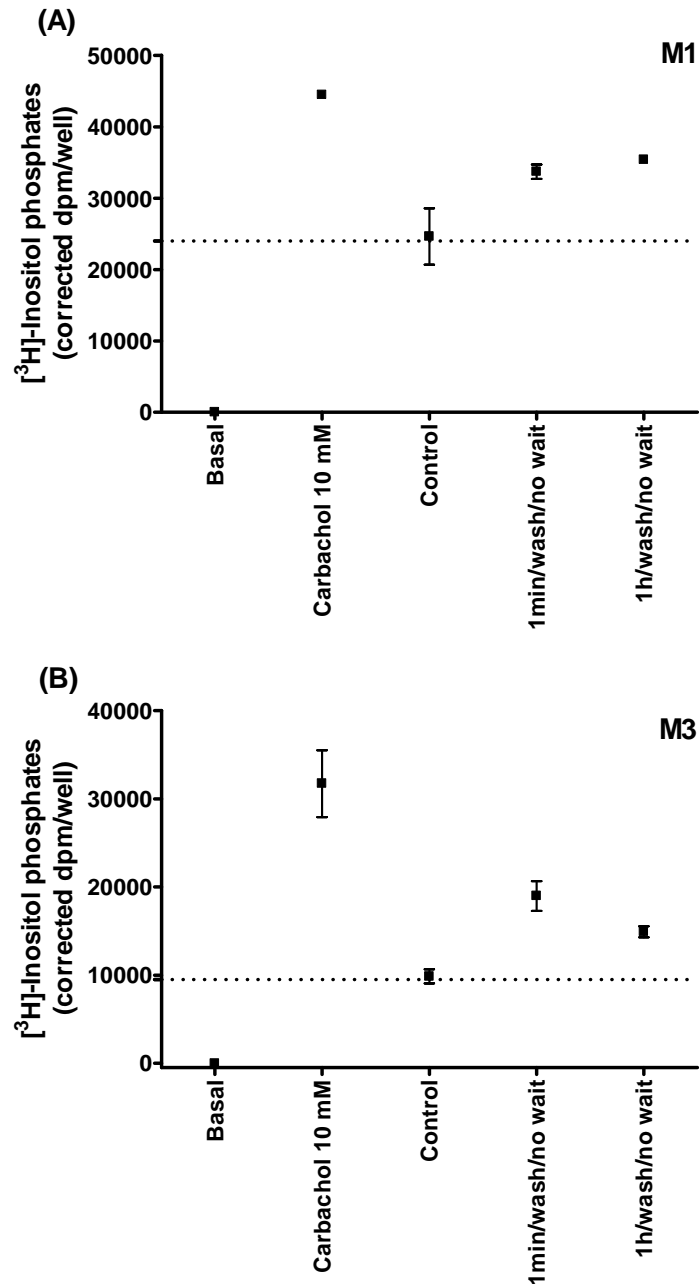
**Figure 4. Effects of acute xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM1 receptor.** Cells were pretreated with 300 nM xanomeline for 1 minute (closed circles) or 1 hour (closed triangles) followed by washing and immediate use. A) Untreated (closed squares) or xanomeline pretreated cells were incubated with increasing concentrations of carbachol in monolayer for 1 hour at 37°C in the presence of 10 mM lithium chloride to determine inositol phosphate production. Alternatively, cells in suspension were treated with increasing concentrations of B) carbachol, C) pilocarpine, or D) xanomeline and the accumulation of inositol phosphates was determined over a 1-hour incubation period at 37°C in the presence of 10 mM lithium chloride for untreated cells (closed squares) and those pretreated with xanomeline. Values represent the means  $\pm$  S.E. of 3-9 experiments conducted in triplicate. Data from (C) and (D) provided by Kayla De Lorme.

Figure 5



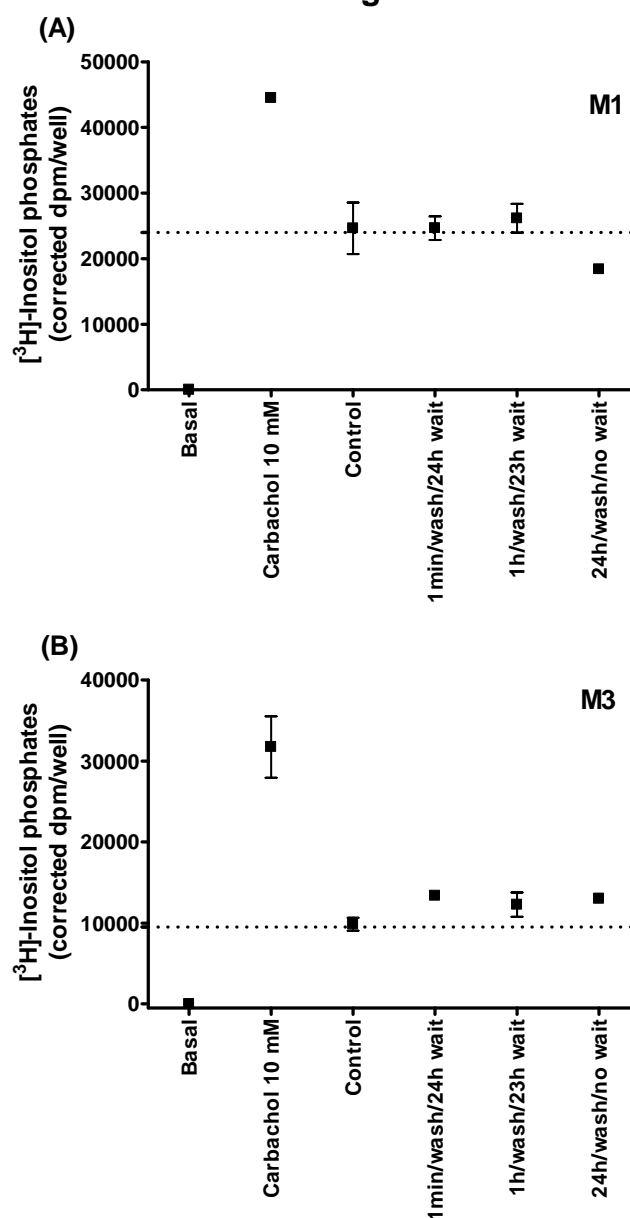
**Figure 5. Effects of acute pilocarpine on the ability of pilocarpine to subsequently stimulate the production of inositol phosphates at the hM3 receptor.** Cells expressing the M3 receptor were pretreated for 1 hour with 10  $\mu$ M pilocarpine followed by washing (closed triangles). Subsequently, untreated (closed squares) or pretreated cells were stimulated with increasing concentrations of pilocarpine for 1 hour at 37°C in the presence of 10 mM lithium chloride to determine the production of inositol phosphates. Values represent the mean  $\pm$  S.E. of 3 experiments conducted in triplicate.

Figure 6



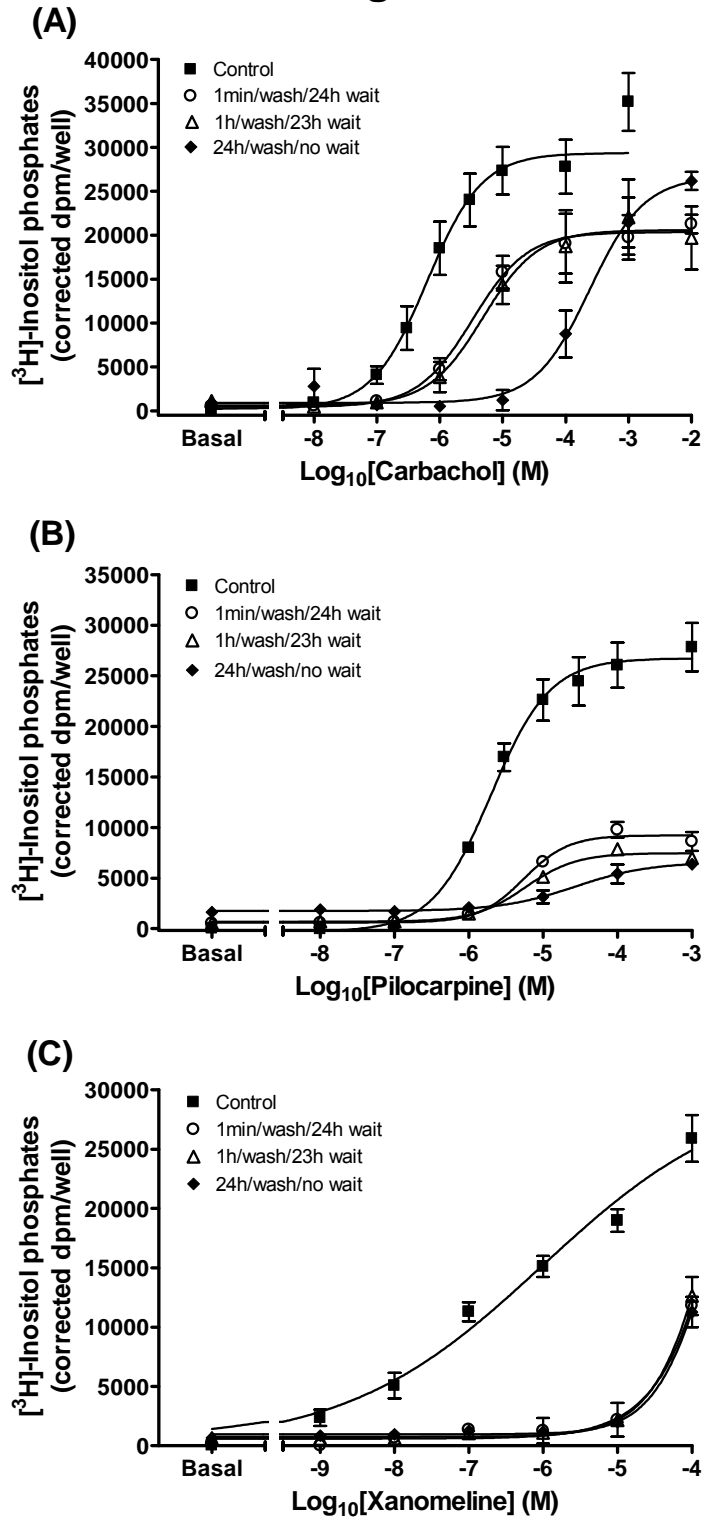
**Figure 6. Stimulation of inositol phosphate production by exogenous phospholipase C following brief xanomeline pretreatments in cells expressing M1 and hM3 receptors.** Cells were pretreated with xanomeline for 1 minute or 1 hour followed by washing. Subsequently, untreated or pretreated cells were stimulated with 2.5 U of phospholipase C for 1 hour at 37°C in the presence of 10 mM lithium chloride and the level of inositol phosphate production was determined. As a reference untreated cells were unstimulated or stimulated with 10  $\mu\text{M}$  carbachol instead of phospholipase C. A) M1 receptor and B) M3 receptor. Values represent the mean of 2 experiments conducted in triplicate.

Figure 7



**Figure 7. Stimulation of inositol phosphate production by exogenous phospholipase C following long-term xanomeline pretreatments in cells expressing hM1 and hM3 receptors.** Cells were pretreated with xanomeline for 1 minute or 1 hour followed by washing and incubation in control media for 24 or 23 hours, respectively. Additional cells were treated with xanomeline for 24 hours prior to washing. Subsequently, untreated or pretreated cells were stimulated with 2.5 U of phospholipase C for 1 hour at 37°C in the presence of 10 mM lithium chloride and the production of inositol phosphates was determined. As a reference untreated cells were unstimulated or stimulated with 10  $\mu\text{M}$  carbachol instead of phospholipase C. A) M1 receptor and B) M3 receptor. Values represent the mean of 2 experiments conducted in triplicate.

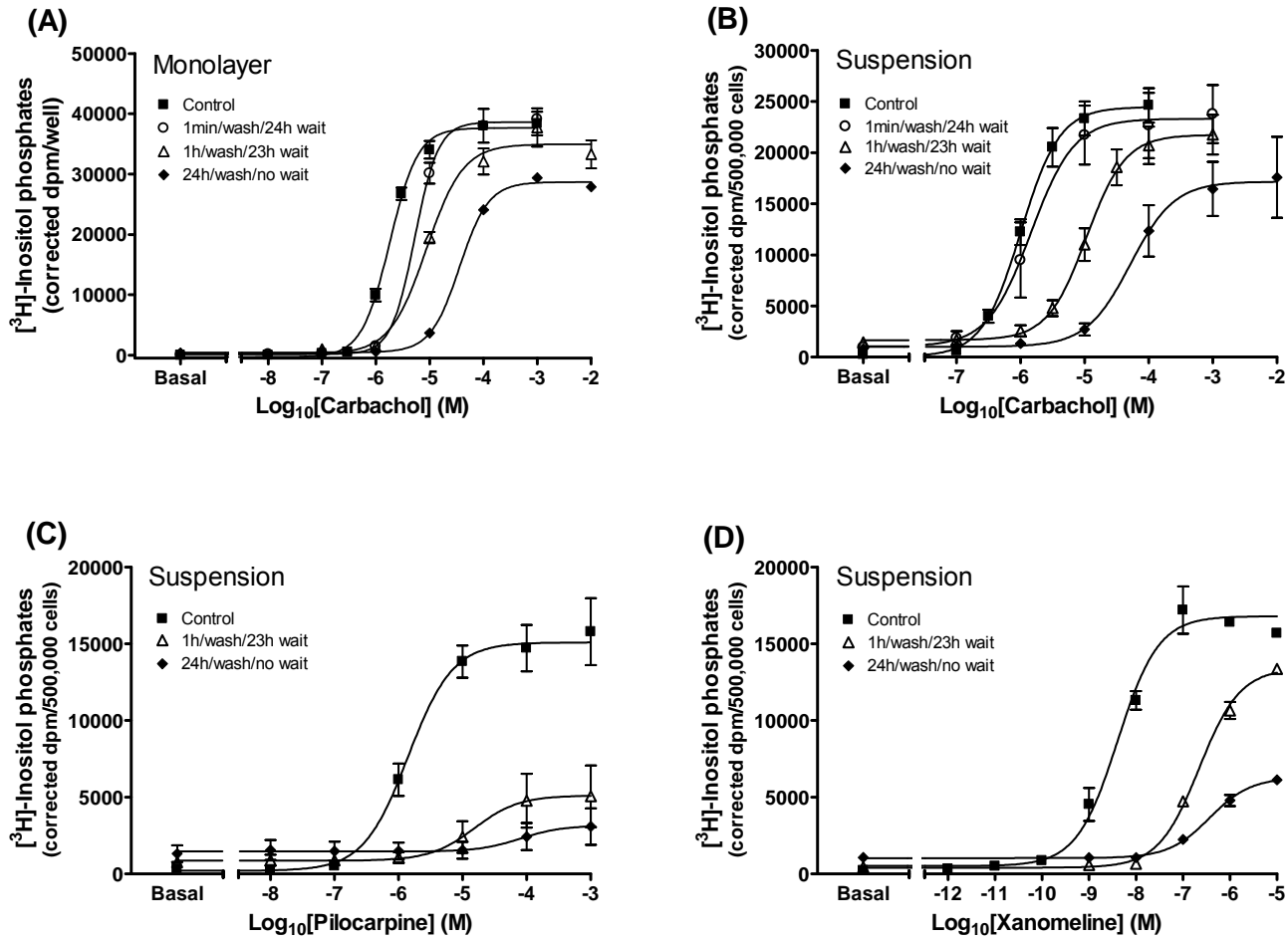
Figure 8





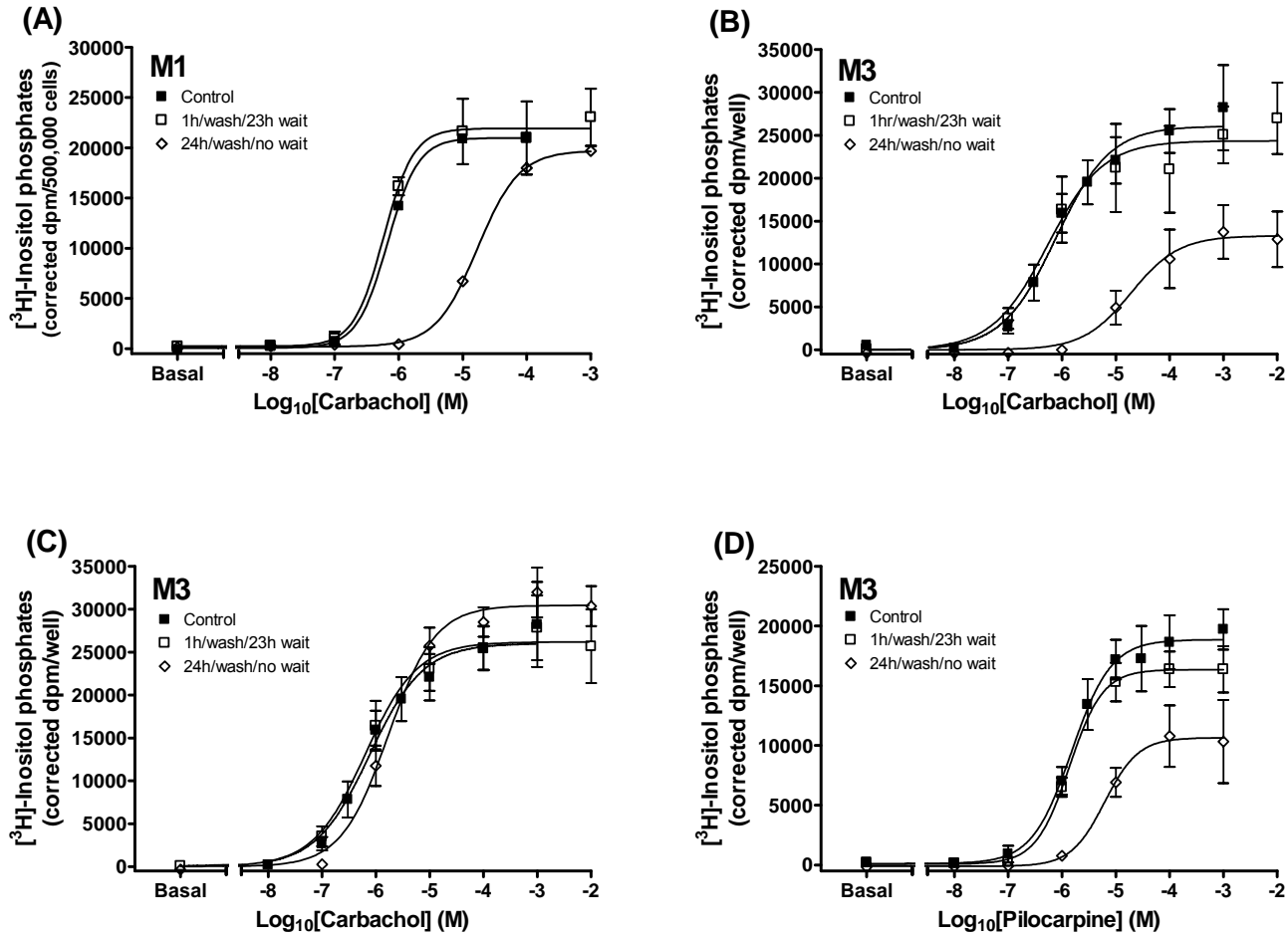
**Figure 8. Effects of prolonged xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM3 receptor.** Cells were pretreated with 10  $\mu$ M xanomeline for 1 minute (open circles) or 1 hour (open triangles) followed by washing and waiting 24 or 23 hours, respectively. An additional group of cells was treated with xanomeline for 24 hours (closed diamonds) prior to washing. Subsequently, cells were treated with increasing concentrations of A) carbachol, B) pilocarpine, or C) xanomeline and the accumulation of inositol phosphates was determined over a 1-hour incubation period at 37°C in the presence of 10 mM lithium chloride for untreated cells (closed squares) or those pretreated with xanomeline. Values represent the means  $\pm$  S.E. of 3-7 experiments conducted in triplicate.

Figure 9



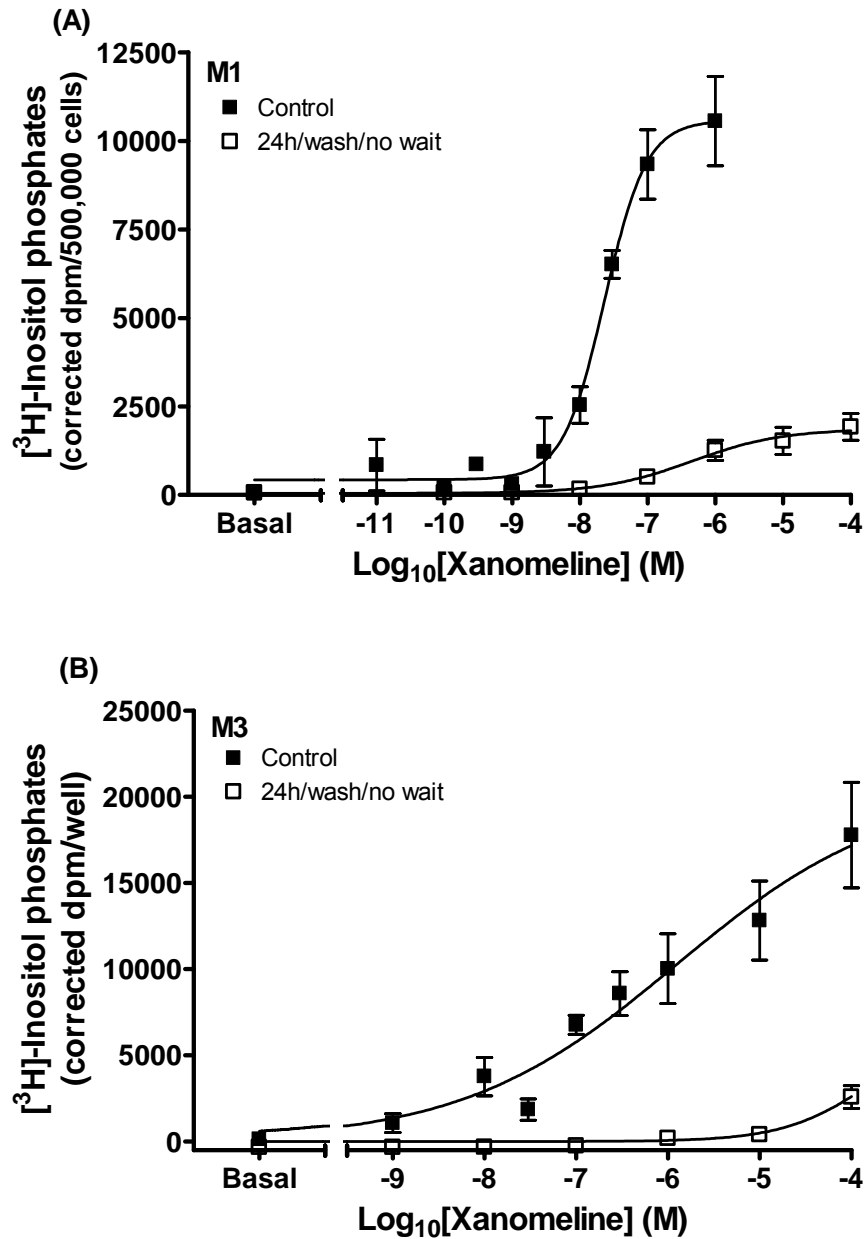
**Figure 9. Effects of prolonged xanomeline exposure on the ability of muscarinic agonists to stimulate the production of inositol phosphates at the hM1 receptor.** Cells were pretreated with 300 nM xanomeline for 1 minute (open circles) or 1 hour (open triangles) followed by washing and waiting 24 or 23 hours, respectively. An additional group of cells was treated with xanomeline for 24 hours (closed diamonds) prior to washing. A) Untreated (closed squares) or xanomeline pretreated cells were incubated with increasing concentrations of carbachol in monolayer for 1 hour at 37°C in the presence of 10 mM lithium chloride to determine inositol phosphate production. Alternatively, cells were treated with increasing concentrations of B) carbachol, C) pilocarpine, or D) xanomeline in suspension and the accumulation of inositol phosphates was determined over a 1-hour incubation period at 37°C in the presence of 10 mM lithium chloride for untreated cells (closed squares) or those pretreated with xanomeline. Values represent the means  $\pm$  S.E. of 2-9 experiments conducted in triplicate. Data from (C) and (D) provided by Kayla De Lorme.

Figure 10



**Figure 10. Effects of long-term agonist exposure on the ability of various muscarinic agonists to stimulate inositol phosphates production at the hM1 and hM3 receptors.** A) Cells expressing the M1 receptor were treated with 10  $\mu$ M carbachol for 1 hour followed by washing and waiting 23 hours in control media (open squares) or treated for 24 hours with carbachol prior to washing (open diamonds). Subsequently, cells were stimulated in suspension with increasing concentrations of carbachol for 1 hour at 37°C in the presence of 10 mM lithium chloride to determine inositol phosphate production. B) Cells expressing the M3 receptor were pretreated with 10  $\mu$ M carbachol as in (A). Subsequently, cells were stimulated in monolayer with increasing concentrations of carbachol and the accumulation of inositol phosphates was determined in the presence of 10 mM LiCl over a 1 hour incubation period at 37°C for untreated cells (closed squares) and those pretreated with carbachol. C) Cells expressing the M3 receptor were treated with 10  $\mu$ M pilocarpine similar to (B) then stimulated with carbachol. D) Cells expressing the M3 receptor were treated with 10  $\mu$ M pilocarpine as in (B) then stimulated with pilocarpine. Values represent the means  $\pm$  S.E. of 2-6 experiments conducted in triplicate.

Figure 11



**Figure 11. Effects of long-term exposure to carbachol on the ability of xanomeline to stimulate inositol phosphate production at hM1 and hM3 receptors.** Cells were pretreated with 10  $\mu$ M carbachol for 24 hours prior to washing (open squares). Subsequently, untreated (closed squares) or pretreated cells were stimulated with increasing concentrations of xanomeline for 1 hour at 37°C in the presence of 10 mM lithium chloride and the production of inositol phosphates was determined. A) Experiments were conducted at the M1 receptor in suspension. B) Experiments were conducted at the M3 receptor in monolayer. Values represent the means of  $\pm$  S.E. of 3 experiments conducted in triplicate. Data in (A) provided by Marianne Grant.

**Table 1. Effects of agonist treatment on PI hydrolysis in CHO cells expressing hM1 and hM3 receptor subtypes.** Cells were incubated at 37°C for 1 hour with increasing concentrations of carbachol (10 nM – 100 µM), pilocarpine (100 nM – 4.5 µM) or xanomeline (1 nM – 10 µM) and the levels of inositol phosphate production were determined in the presence of 10 mM lithium chloride. Additional groups of cells were treated with xanomeline (1 nM – 100 µM) for 1 minute or 1 hour followed by washing and use in the functional assay. Parameters ( $\pm$  S.E.M.) were derived from computer-assisted non-linear regression analysis.

	<b>M1</b>		<b>M3</b>	
	<b>pEC<sub>50</sub><sup>a</sup></b>	<b>E<sub>max</sub><sup>b</sup></b>	<b>pEC<sub>50</sub></b>	<b>E<sub>max</sub></b>
<b>Agonist stimulation</b>				
Carbachol	5.8 $\pm$ 0.24	15800 $\pm$ 2600	6.3 $\pm$ 0.04	23300 $\pm$ 3900
Pilocarpine			5.8 $\pm$ 0.08	16400 $\pm$ 2100
Xanomeline	8.0 $\pm$ 0.11 <sup>†</sup>	14700 $\pm$ 1400	6.9 $\pm$ 0.07 <sup>†</sup>	11100 $\pm$ 1300*
<b>Xanomeline pretreatment</b>				
1 min/washout/no wait	7.1 $\pm$ 0.08 <sup>§</sup>	11100 $\pm$ 350	6.1 $\pm$ 0.10	9800 $\pm$ 2000
1 h/washout/no wait	7.6 $\pm$ 0.11	10200 $\pm$ 1200	7.1 $\pm$ 0.52	6200 $\pm$ 1300

<sup>a</sup> Negative logarithm of the midpoint (potency) parameter.

<sup>b</sup> Maximal response. Values are expressed as dpm/well after correcting for standard recovery.

No statistical difference in E<sub>max</sub> was detected at the M1 receptor between xanomeline and carbachol.

<sup>§</sup> ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in EC<sub>50</sub> where indicated between xanomeline 1min/wash/no wait or 1h/wash/no wait compared with xanomeline at the M1 receptor.

<sup>†</sup> ANOVA followed by Tukey's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> between xanomeline compared with carbachol or pilocarpine at the M1 or M3 receptor.

\* ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in E<sub>max</sub> between xanomeline compared with carbachol at the M3 receptor.

No statistical differences in E<sub>max</sub> or pEC<sub>50</sub> were found between the xanomeline-pretreated groups compared to xanomeline at the M3 receptor.

**Table 2. Effects of carbachol, pilocarpine or xanomeline stimulation following short- and long-term agonist pretreatment and washout on inositol phosphates production in CHO cells expressing hM3 receptors.** Cells were pretreated with 10  $\mu$ M agonists for 1 minute or 1 hour followed by washing and immediate use in the functional assay or subsequent to incubation for 24 or 23 hours, respectively, in the absence of free agonist. Additional groups of cells were treated with 10  $\mu$ M agonist for 24 hours prior to washing. In the functional assay, cells were stimulated with increasing concentrations of carbachol, pilocarpine or xanomeline for 1 hour at 37°C in the presence of 10 mM lithium chloride. Parameters ( $\pm$  S.E.M.) were derived from computer-assisted non-linear regression analysis.

	<u>Carbachol Stimulation</u>		<u>Pilocarpine Stimulation</u>		<u>Xanomeline Stimulation</u>	
	pEC <sub>50</sub> <sup>a</sup>	E <sub>max</sub> <sup>b</sup>	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>
<b>Xanomeline</b>						
Control (untreated)	6.2 $\pm$ 0.09	28300 $\pm$ 3100	5.7 $\pm$ 0.004	26800 $\pm$ 2300	6.2 $\pm$ 0.14	28100 $\pm$ 3000
1 min/washout/no wait	4.7 $\pm$ 0.19*	30700 $\pm$ 4100	4.9 $\pm$ 0.05 <sup>†</sup>	24500 $\pm$ 2800	4.5 $\pm$ 0.10 <sup>‡</sup>	32200 $\pm$ 2800
1 h/washout/no wait	4.2 $\pm$ 0.06*	28900 $\pm$ 5100	4.3 $\pm$ 0.05 <sup>†</sup>	17900 $\pm$ 1500 <sup>†</sup>	4.4 $\pm$ 0.15 <sup>‡</sup>	32200 $\pm$ 2800
Xanomeline & carbachol – co-addition	3.7 $\pm$ 0.11*	24000 $\pm$ 4800				
1 min/washout/24h wait	5.5 $\pm$ 0.14*	20400 $\pm$ 2000	5.3 $\pm$ 0.06 <sup>†</sup>	9200 $\pm$ 880 <sup>†</sup>	ND	ND
1 h/washout/23h wait	5.1 $\pm$ 0.10*	25600 $\pm$ 2700	5.2 $\pm$ 0.05 <sup>†</sup>	7000 $\pm$ 600 <sup>†</sup>	ND	ND
24 h/washout/no wait	3.6 $\pm$ 0.20*	26800 $\pm$ 1000	4.0 $\pm$ 0.04 <sup>†</sup>	6600 $\pm$ 500 <sup>†</sup>	ND	ND
<b>Carbachol</b>						
Control (untreated)	6.1 $\pm$ 0.11	25400 $\pm$ 2400			5.7 $\pm$ 0.35	21300 $\pm$ 3300
1 h/washout/23 h wait	6.0 $\pm$ 0.43	25600 $\pm$ 3300				
24 h/washout/no wait	4.7 $\pm$ 0.27 <sup>s</sup>	13200 $\pm$ 3200 <sup>s</sup>			ND	ND



Table 2 cont.

	<u>Carbachol Stimulation</u>		<u>Pilocarpine Stimulation</u>		<u>Xanomeline Stimulation</u>	
	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>
<b>Pilocarpine</b>						
Control (untreated)	6.1 ± 0.11	25400 ± 2400	5.8 ± 0.04	19000 ± 1900		
1 h/washout/23 h wait	6.2 ± 0.08	26300 ± 3400	5.9 ± 0.03	16400 ± 1600		
24 h/washout/no wait	5.8 ± 0.11	30500 ± 2300	5.3 ± 0.11 <sup>#</sup>	10800 ± 3200		

<sup>a</sup> Negative logarithm of the midpoint (potency) parameter.

<sup>b</sup> Maximal response. Values are expressed as dpm/well after correcting for standard recovery.

ND – parameter value was not determined

\* ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with carbachol stimulation compared to carbachol stimulation in untreated cells.

† ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with pilocarpine stimulation compared to pilocarpine stimulation in untreated cells.

‡ ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with xanomeline stimulation compared to xanomeline stimulation in untreated cells.

§ ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between carbachol-pretreated groups with carbachol stimulation compared to carbachol stimulation in untreated cells.

# ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between pilocarpine-pretreated groups with pilocarpine compared to pilocarpine stimulation in untreated cells.

No significant difference was detected between pilocarpine-pretreated groups with carbachol stimulation compared to carbachol stimulation in untreated cells.

**Table 3. Effects of carbachol, pilocarpine or xanomeline stimulation following short- and long-term agonist pretreatment and washout on inositol phosphates production in CHO cells expressing hM1 receptors.** Cells were pretreated with 300 nM xanomeline or 10  $\mu$ M carbachol for 1 minute or 1 hour followed by washing and immediate use in the functional assay or subsequent to incubation for 24 or 23 hours, respectively, in the absence of free agonist. Additional groups of cells were treated with 300 nM xanomeline or 10  $\mu$ M carbachol for 24 hours prior to washing. In the functional assay cells, were stimulated with increasing concentrations of carbachol, pilocarpine or xanomeline for 1 hour at 37°C in the presence of 10 mM lithium chloride. Parameters ( $\pm$  S.E.M.) were derived from computer-assisted non-linear regression analysis.

Pretreatment	<u>Carbachol Stimulation</u>		<u>Pilocarpine Stimulation</u>		<u>Xanomeline Stimulation</u>	
	pEC <sub>50</sub> <sup>a</sup>	E <sub>max</sub> <sup>b</sup>	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>
<b>Xanomeline – suspension<sup>c</sup></b>						
Control (untreated)	6.0 $\pm$ 0.06	24500 $\pm$ 1800	5.8 $\pm$ 0.12	13900 $\pm$ 1700	7.9 $\pm$ 0.40	18200 $\pm$ 1200
1 min/washout/no wait	6.4 $\pm$ 0.18*	25100 $\pm$ 2200				
1 h/washout/no wait	ND	21000 $\pm$ 2100	5.6 $\pm$ 0.75	12800 $\pm$ 1000	6.4 $\pm$ 0.55 <sup>§</sup>	17900 $\pm$ 1200
1 min/washout/24 h wait	5.9 $\pm$ 0.10	23300 $\pm$ 2900				
1 h/washout/23 h wait	5.0 $\pm$ 0.06*	21100 $\pm$ 2400	4.3 $\pm$ .029 <sup>†</sup>	6300 $\pm$ 1800 <sup>†</sup>	6.6 $\pm$ 0.03	11600 $\pm$ 2200 <sup>§</sup>
24 h/washout/no wait	4.3 $\pm$ 0.11*	14700 $\pm$ 2800*	3.9 $\pm$ 0.29 <sup>†</sup>	3800 $\pm$ 960 <sup>†</sup>	6.3 $\pm$ 0.09 <sup>§</sup>	5100 $\pm$ 1200 <sup>§</sup>
<b>Xanomeline – monolayer<sup>d</sup></b>						
Control (untreated)	5.7	37700				
1 min/washout/no wait	5.6	35500				
1 h/washout/no wait	5.5	31500				
1 min/washout/24 h wait	5.3	38600				
1 h/washout/23 h wait	5.1	35000				
24 h/washout/no wait	4.4	28700				

Table 3 cont.

	<u>Carbachol Stimulation</u>		<u>Pilocarpine Stimulation</u>		<u>Xanomeline Stimulation</u>	
	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>
<b>Carbachol</b>						
Control (untreated)	6.2	21000			7.7 ± 0.11	10900 ± 1100
1 h/washout/23 h wait	6.2 ± 0.15	21900 ± 3300				
24 h/washout/no wait	4.8	19700			6.6 ± 0.01 <sup>‡</sup>	1600 ± 370 <sup>‡</sup>

<sup>a</sup> Negative logarithm of the midpoint (potency) parameter.

<sup>b</sup> Maximal response. Values are expressed as dpm/well after correcting for standard recovery.

<sup>c</sup> Pretreatments were conducted in monolayer and the functional assay was conducted in suspension.

<sup>d</sup> Pretreatments and the functional assay were conducted on cells in monolayer. The values listed are for a single assay conducted in triplicate.

ND – parameter not determined

\* ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with carbachol stimulation compared to carbachol stimulation in untreated cells.

† ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with pilocarpine stimulation compared to pilocarpine stimulation in untreated cells.

§ ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> where indicated between xanomeline-pretreated groups with xanomeline stimulation compared to xanomeline stimulation in untreated cells.

‡ Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pEC<sub>50</sub> and E<sub>max</sub> between carbachol 24h/wash/no wait with xanomeline stimulation compared to xanomeline stimulation in untreated cells.

**CHAPTER 5: Receptor activation is required for the development of the long-term effects of xanomeline at the M1 and M3 muscarinic receptors**

## **Introduction**

Binding of conventional ligands to muscarinic receptors occurs at the orthosteric site of the receptor. This binding results in activation of down-stream signaling cascades through conformational changes in transmembrane domains 3, 5, 6 and 7 (Eglen, 2005). The M1 and M3 receptors are coupled to G<sub>q/11</sub> G proteins that lead to the activation of the phospholipase C signaling pathway (Volpicelli & Levey, 2004). Prolonged activation of the receptor through the orthosteric binding site can lead to changes in receptor expression and a decrease in the ability of subsequent agonist treatment to stimulate a functional response (Hu et al., 1991; Maloteaux & Hermans, 1994). Additionally, binding of a drug to an allosteric site on a receptor may lead to changes in the conformation of the receptor's orthosteric domain that can result in receptor activation (Christopoulos & Kenakin, 2002; Wess, 2005). Examples of such allosteric muscarinic receptor agonists include AC-42 (Spalding et al., 2002) and WIN51708 (Lanzafame et al., 2006).

Xanomeline has been shown to bind in a wash-resistant manner at the M1 receptor (Christopoulos et al., 1998; Jakubik et al., 2002). Further results demonstrated that binding occurs in this manner even when the primary binding site is blocked (Jakubik et al., 2002). When the orthosteric site is blocked acutely with atropine, stimulation of the receptor is inhibited; however, when atropine is washed out xanomeline is again able to stimulate a functional response (Christopoulos et al., 1998). Xanomeline has been shown to bind in an allosteric manner; however, it is unclear if activation through the orthosteric site is necessary for the development of the long-term consequences of this binding.

Thus, the aim of the study was to determine if the orthosteric site is necessary for the long-term consequences of xanomeline binding at the M1 and M3 receptors. To determine this, we used a variety of protocols to block receptor function in CHO cells

expressing either M1 or M3 muscarinic receptors. Our data demonstrate that receptor activation through the orthosteric site is necessary for the long-term consequence of xanomeline on receptor binding and functional properties. However, xanomeline binding in the absence of function is still able to regulate the affinity of the receptor for other ligands.

## **Results**

### **Effects of atropine on long-term xanomeline-induced inhibition of [<sup>3</sup>H]NMS binding at the M1 and M3 receptors**

Experiments were designed to determine the role of xanomeline interaction with the receptor's orthosteric domain in the development of its long-term effects. A receptor-saturating concentration of atropine (10  $\mu$ M), a muscarinic receptor antagonist, was added either concurrently with 10  $\mu$ M xanomeline pretreatment or during the waiting period following xanomeline washout. [<sup>3</sup>H]NMS saturation binding isotherms were constructed to examine the effects of blocking the orthosteric site with atropine on xanomeline-induced changes in both radioligand affinity and maximal binding at the M3 receptor. The simultaneous presence of atropine during incubation of cells with xanomeline for one hour followed by washing and waiting 23 hours in control medium did not influence the significant ( $p < 0.05$ ) decrease in [<sup>3</sup>H]NMS maximal binding induced by xanomeline pretreatment alone (Fig. 1 a, b; Table 1). However, when atropine was present during the 23 hour incubation following xanomeline pretreatment and washing, the long-term changes in radioligand maximal binding were no longer evident (Fig. 1 c; Table 1). In contrast, either protocol of atropine treatment abolished the long-term effects of xanomeline on radioligand affinity. All treatment groups had similar protein content.

When parallel experiments were conducted at the M1 receptor, similar results were seen to those obtained at the M3 receptor. When atropine (10  $\mu$ M) and xanomeline (3  $\mu$ M) were simultaneously present for one hour followed by washing and waiting, the significant ( $p < 0.05$ ) decrease in maximal binding was not affected compared to xanomeline pretreatment alone (Fig. 2 a, b). In contrast, when cells were treated with atropine for 23 hours subsequent to one hour xanomeline pretreatment the long-term changes in maximal binding were no longer evident (Fig. 2 c). Additionally, in both cases atropine treatment prevented the change in radioligand binding affinity observed with xanomeline treatment alone (Fig. 2 b, c). Results are summarized in Table 1.

#### **Effects of atropine on xanomeline-induced long-term alterations in receptor sensitivity at the M1 and M3 receptors**

Similar to the previously described [ $^3$ H]NMS binding experiments, functional experiments were conducted to determine the effects of blocking the primary binding site with a receptor-saturating concentration of atropine (10  $\mu$ M), either concurrently with 10  $\mu$ M xanomeline pretreatment for one hour or during the 23 hour waiting period following xanomeline pretreatment and washing on the ability of increasing concentrations of carbachol to stimulate the production of inositol phosphates in cells expressing the M3 receptor. In contrast to results obtained from binding experiments, simultaneous presence of atropine with xanomeline for one hour followed by washing and waiting for 23 hours in control medium abolished the effects of xanomeline on carbachol potency (Fig. 3 a, b; Table 1). The presence of atropine during the 23 hour incubation following xanomeline treatment and washout also eliminated xanomeline-induced changes in

carbachol potency, although a decrease in efficacy was observed compared to control (Fig. 3 a, c; Table 1).

Functional experiments were also employed using atropine (10  $\mu$ M) to block the orthosteric site of the M1 receptor to determine the effects of blocking receptor function on the ability of increasing concentrations of carbachol to stimulate inositol phosphate production subsequent to long-term exposure to xanomeline (300 nM). When cells were treated concurrently with atropine and xanomeline for one hour followed by washing and waiting 23 hours, the effect on carbachol potency was no longer evident compared to xanomeline pretreatment alone (Fig. 4 a, b), similar to the observed effects at the M3 receptor. However, the basal level of receptor stimulation was elevated at the M1 receptor under these treatment conditions, which was not observed with xanomeline treatment alone (Fig. 4 b). The reason for the increase in basal stimulation is unclear. Similar to the results obtained with saturation binding, the presence of atropine during the 23 hour waiting period subsequent to xanomeline pretreatment abolished the changes observed with xanomeline treatment alone (Fig. 4 c). Additionally, no change in maximal stimulation was evident with either atropine pretreatment condition. Results are summarized in Table 1.

#### **Effects of blocking receptor function, using a non-functional mutant, on inhibition of [<sup>3</sup>H]NMS and saturation binding parameters at the M1 receptor**

Experiments were designed to further test the necessity of receptor activation on the long-term effects of xanomeline binding at the M1 receptor using a non-functional receptor mutant (rM1<sup>123</sup>) expressed in CHO cells. In this case, the mutant receptor was cloned from rat so additional parallel experiments were conducted with the rat wild-type receptor (rM1). Lack of a functional response of the receptor mutant was confirmed



under our experimental conditions (Fig. 5 a). Exposure of wild-type rM1 receptors to carbachol or xanomeline indicated that xanomeline is a partial agonist in stimulating inositol phosphate production. This is in contrast to the hM1 receptor where xanomeline and carbachol had similar efficacies (Fig. 5 a; Chapter 4 Fig. 2 a). However, at either wild-type receptor, human or rat, xanomeline was significantly ( $p < 0.05$ ) more potent than carbachol (Fig. 5 a; Chapter 4 Fig. 2 a). Subsequently, inhibition of [<sup>3</sup>H]NMS binding experiments were conducted to compare the ability of xanomeline to regulate the rM1 and rM1<sup>123</sup> mutant receptors.

When cells expressing either the rM1 or rM1<sup>123</sup> receptors were exposed to increasing concentrations of xanomeline in the binding assay, the potencies and maximal inhibition of [<sup>3</sup>H]NMS binding were similar between receptors (Fig. 5 b, c; Table 2). Similarly, when cells were exposed to xanomeline for one hour followed by washing and used immediately in the binding assay the potencies and maximal inhibition of [<sup>3</sup>H]NMS binding were nearly identical between the receptors. However, the potency decreased by approximately 1.3 – 1.6 orders of magnitude compared to the former treatment. In contrast, when cells were exposed to xanomeline for one hour followed by washing and waiting 23 hours a biphasic binding curve was observed at the rM1 receptor, but a monophasic binding curve was observed at the rM1<sup>123</sup> mutant receptor (Fig. 5 b, c). In this case the potency at the low-potency site for the rM1 receptor was similar to the potency at the rM1<sup>123</sup> mutant receptor. Similar results were observed in the binding isotherms following 24 hour xanomeline treatment prior to washing, with a biphasic binding curve observed at the rM1 and a monophasic binding curve at the rM1<sup>123</sup> mutant receptor. Interestingly, the potency for 24 hour xanomeline treatment followed by washing was indistinguishable from the potency of the acute xanomeline treatment at the rM1<sup>123</sup> mutant receptor (Fig. 5 c). In the case of long-term exposure to

xanomeline, inhibition of [<sup>3</sup>H]NMS binding was complete for all groups except xanomeline treatment for one hour followed by washing and further incubation in control media for 23 hours. In that case the binding curve was still decreasing at the highest concentrations of xanomeline tested, but the vehicle that xanomeline was dissolved in (DMSO) prohibited testing at higher concentrations. Results are summarized in Table 2.

Additionally, saturation binding experiments were conducted at the rM1 wild-type and rM1<sup>123</sup> mutant receptors to further determine the importance of receptor function on cell-surface receptor density and radioligand binding affinity. In both cases, when cells were exposed to xanomeline (3 μM) for one hour followed by washing and immediate use there was no change in cell-surface receptor density, but a decrease in radioligand affinity compared to their respective controls (Fig. 5 d, e). When rM1 expressing cells were treated with xanomeline for one hour followed by washing and waiting 23 hours there was a significant ( $p < 0.05$ ) decrease in cell-surface receptor density with no change in radioligand affinity compared to control (Fig. 5 d). In contrast, the decrease in cell-surface receptor density compared to control was absent in cells expressing the rM1<sup>123</sup> mutant receptor (Fig. 5 e). Similarly, when cells expressing the wild-type rM1 receptor were exposed to xanomeline for 24 hours prior to washing there was a decrease in cell-surface receptor density along with a significant ( $p < 0.05$ ) decrease in radioligand affinity (Fig. 5 d). In contrast, the decrease in cell-surface receptor density was absent and the decrease in radioligand affinity was less pronounced in cells expressing the rM1<sup>123</sup> mutant receptor (Fig. 5 e). Results are summarized in Table 2.

Due to the fact that xanomeline was dissolved in dimethyl sulfoxide (DMSO), control experiments were designed to ensure that DMSO did not contribute to the observed effects of xanomeline. Cells were treated with DMSO concentrations corresponding to those in contact with cells during the various treatment modalities with

xanomeline in the inhibition of [<sup>3</sup>H]NMS binding assay. DMSO did not affect radioligand binding at the highest concentrations used to prepare xanomeline dilutions (1%; Fig. 6 a, b) in either case. Additionally, protein content was determined following DMSO pretreatments for the various conditions. Once again, there was no change in protein content for any of the treatment groups (Fig. 6 c, d).

As previously mentioned, there was a decrease in protein expression following xanomeline treatments at the M1 receptor, thus protein analysis was conducted using the various xanomeline pretreatments previously described. In this case there was no difference in protein content for any of the aforementioned xanomeline pretreatments (Fig. 6 e, f).

### **Effects of blocking receptor function using siRNA targeted against G<sub>q</sub> and G<sub>11</sub> G proteins at the M3 receptor**

Experiments were conducted to further examine the role of receptor activation on the long-term effects of xanomeline treatments on [<sup>3</sup>H]NMS saturation binding parameters at the M3 receptor. It is known that the M3 receptor exerts its functional response via activation of the G<sub>q/11</sub> class of G proteins (Volpicelli & Levey, 2004). Thus, we utilized siRNA technology to block expression of these G proteins. To this effect, cells were transfected with 50 nM siRNA targeting G<sub>q</sub> and G<sub>11</sub> G proteins in an effort to block receptor function. Western blot analysis confirmed that transfection with siRNA targeting G<sub>q/11</sub> G proteins resulted in a nearly 100 percent decrease in the detectable expression levels of G<sub>q/11</sub> class of G proteins (Fig. 7 a). In addition, loss of receptor function following transfection was assessed by determining agonist-mediated production of inositol phosphates using two concentrations of carbachol (1 μM and 100 μM) or xanomeline (100 nM and 10 μM). The response to carbachol was decreased by

62 and 27 percent, respectively, and the response to xanomeline was reduced by 78 and 82 percent, respectively (Fig. 7 b). The ability of agonist to stimulate a functional response subsequent to siRNA treatment suggests that although the levels of G<sub>q/11</sub> G proteins were no longer detectable following siRNA treatment some G<sub>q/11</sub> G proteins were still present. A [<sup>3</sup>H]NMS saturation binding paradigm was once again applied to determine the effects of blocking receptor function on xanomeline-induced changes in both radioligand affinity and maximal binding. However, while attached cells were pretreated with xanomeline, binding experiments were conducted in suspension due to technical limitations (see Methods). In control non-siRNA transfected cells, a trend toward a decrease in radioligand affinity was observed following one hour, wash and immediate use xanomeline pretreatment as well as pretreatment with xanomeline for 24 hours prior to washing (Fig. 7 c; Table 3). Additionally, a trend toward a decrease in maximal binding resulted following one hour xanomeline pretreatment followed by washing and waiting 23 hours or for 24 hour xanomeline treatment followed by washing and no waiting (Fig. 7 c; Table 3). These effects on radioligand affinity and maximal binding were similar, although less pronounced, than those obtained from [<sup>3</sup>H]NMS saturation binding performed in monolayer (Chapter 3 Fig. 19; Chapter 3 Table 2). This is similar to previous work by Thompson & Fisher (1990) in which there was a greater loss of receptors in cells that were pretreated with carbachol in monolayer versus those in cells pretreated in suspension prior to the radioligand binding assay. Cells transfected with siRNA displayed a trend toward a decrease in radioligand affinity for all xanomeline pretreatment groups except the 24 hour treatment group which had a significant ( $p < 0.05$ ) decrease in radioligand affinity compared to control. There was no change in maximal binding for any of the siRNA-treated groups compared to control (Fig. 7 d; Table 3). Protein content remained the same across the various pretreatment groups.

## **Differential role of wash-resistant binding versus persistent receptor activation in xanomeline long-term effects on saturation binding at the M3 receptor**

Shortening the hydrophobic chain of xanomeline has previously been shown to result in diminishing its wash-resistant binding and functional properties at the M1 receptor (Jakubik et al., 2004). Thus, experiments were conducted at the M3 receptor to determine the ability of xanomeline analogs to bind in a reversible and wash-resistant manner. The analogs studied were hexyl (xanomeline), pentyl (one carbon shorter), propyl (three carbons shorter) and ethyl (four carbons shorter). All analogs tested were able to decrease [<sup>3</sup>H]NMS binding in a concentration dependent manner with complete inhibition when included in the binding assay (Fig. 8 a). The rank order of potencies was pentyl > hexyl ≈ propyl > ethyl. When cells were treated with the various analogs for one hour followed by washing and used immediately there was a concentration dependent decrease in [<sup>3</sup>H]NMS binding for all except the ethyl analog (Fig. 8 b). In this case the rank order of potencies was hexyl ≈ pentyl > propyl. Additionally, the maximal inhibition of [<sup>3</sup>H]NMS binding was greater for the hexyl and pentyl analogs compared to the propyl analog (Table 5). In the assay of receptor function, the propyl analog produced 45% of the PI response to xanomeline; however, it did not display any wash-resistant receptor activation (Fig. 8 c). We therefore utilized the propyl analog to discern the role of persistent receptor activation, versus wash-resistant binding per se in the observed long-term effects of xanomeline at the M3 receptor. When cells were exposed to the propyl analog at 100 μM for one hour followed by washing and overnight incubation in control media, there was only a minute decrease in cell-surface receptor density in conjunction with a slight increase in radioligand affinity (Fig. 8 d). In contrast, 24 hour exposure of cells to the propyl analog prior to washing resulted in an anomalous significant ( $p < 0.05$ )

increase in cell-surface receptor density along with a significant ( $p < 0.05$ ) decrease in the affinity of [<sup>3</sup>H]NMS (Fig. 8 d). Results are summarized in Table 5.

## **Discussion**

While xanomeline wash-resistant binding takes place at an allosteric site on the muscarinic receptor, the persistent functional response likely involves activation of the receptor's primary domain by xanomeline's active head group (Christopoulos et al., 1999; Christopoulos et al., 1998; Jakubik et al., 2002). Thus, further experiments were conducted to determine the role of receptor activation in the long-term changes observed following acute exposure of the receptor to xanomeline (Fig. 1 & Fig. 2). Blocking the primary binding site with atropine during brief treatment with xanomeline did not prevent long-term reductions in receptor availability induced by persistently-bound xanomeline at either the M1 or M3 receptor, suggesting that activation of the receptor during the initial treatment period is not necessary in order for xanomeline to induce long-term changes in either case. These data also indicate that access to the orthosteric site is not necessary for wash-resistant binding of xanomeline to take place. This is in agreement with previous studies at the M1 receptor showing that this interaction takes place at a secondary receptor binding site (Christopoulos et al., 1998; De Lorme et al., 2006; De Lorme et al., 2007). However, blocking the primary binding site with atropine during the long waiting period following washing off free xanomeline blunted the long-term effects of xanomeline on [<sup>3</sup>H]NMS binding at both the M1 and M3 receptor. Together, these results indicate that receptor activation is necessary to produce the long-term consequences of xanomeline wash-resistant binding at both the M1 and M3 receptor.

Functional experiments were conducted to determine the effects of suppressing receptor activation on the delayed actions of xanomeline at M1 and M3 receptors (Fig. 3 & Fig. 4). The concomitant presence of atropine with xanomeline during the initial pretreatment period, followed by washing and prolonged incubation in ligand-free medium, prevented the change in carbachol potency in stimulating PI hydrolysis observed following xanomeline treatment alone at both the M1 and M3 receptors. This is in contrast to the results obtained in [<sup>3</sup>H]NMS saturation binding experiments in which the long-term effects of wash-resistant xanomeline binding were still evident under these conditions. However, at the M1 receptor, the basal level of inositol phosphate production was elevated following simultaneous pretreatment with atropine and xanomeline prior to washing and waiting. The presence of atropine during the waiting period following xanomeline pretreatment prevented the change in carbachol potency observed with xanomeline treatment alone at both subtypes. In comparison, the presence of atropine during the 23 hour incubation following acute xanomeline treatment resulted in a decrease in carbachol efficacy compared to atropine treatment alone at the M3 receptor, which was not evident at the M1 receptor. This is in contrast to the observed antagonism by atropine of xanomeline-induced changes in [<sup>3</sup>H]NMS saturation binding at the M3 receptor. The discrepancies between the results obtained in saturation binding compared to the functional response assays may be due to unique interactions of atropine with the receptor. While commonly described as an antagonist, atropine also exhibits properties of an inverse agonist (Nelson et al., 2006; Thor et al., 2008). For example, previous research found that continuous treatment with atropine can lead to an increase in receptor expression (Lee & Wolfe, 1989; May et al., 2005). Thus, the effects of atropine itself may be concealing the functional effects of long-term xanomeline exposure.

A mutant rM1 receptor was used to further explore the role of receptor activation on the development of the long-term effects of xanomeline at the rM1 receptor. Functional assays were performed to confirm that the mutation resulted in a complete loss of the ability of agonists to stimulate the production of inositol phosphates at the rM1<sup>123</sup> mutant receptor (Fig. 5 a). When cells were exposed acutely to xanomeline it resulted in wash-resistant binding in a concentration dependent manner at both the wild-type and mutant rM1 receptors (Fig. 5 b, c). Interestingly, overnight incubation in control media results in a biphasic binding curve at the wild-type receptor, but a monophasic binding curve for the mutant receptor. Similar results were observed following xanomeline treatment for 24 hours prior to washing. Taken together, these results suggest that receptor function is necessary to develop the prolonged effects of xanomeline binding. The similarity in potency between cells expressing rM1<sup>123</sup> receptors treated with xanomeline for one hour followed by washing and immediate use and those treated with xanomeline for 24 hours prior to washing suggests that persistently bound xanomeline may interfere with [<sup>3</sup>H]NMS binding in the absence of receptor function. To further support this conclusion saturation binding experiments were conducted employing the various xanomeline pretreatments. Under these conditions, prolonged treatment with xanomeline resulted in a decrease in cell-surface receptor density for wild-type, but not rM1<sup>123</sup> mutant receptors (Fig. 5 d, e). Interestingly, 24 hour treatment with xanomeline prior to washing resulted in a decrease in radioligand affinity for both the wild-type and mutant receptors, however, it was more pronounced at the wild-type receptor. Thus, these results demonstrate that receptor function is necessary for the long-term effects of xanomeline on cell-surface receptor density, but receptor function is not solely responsible for the long-term effects on radioligand affinity.



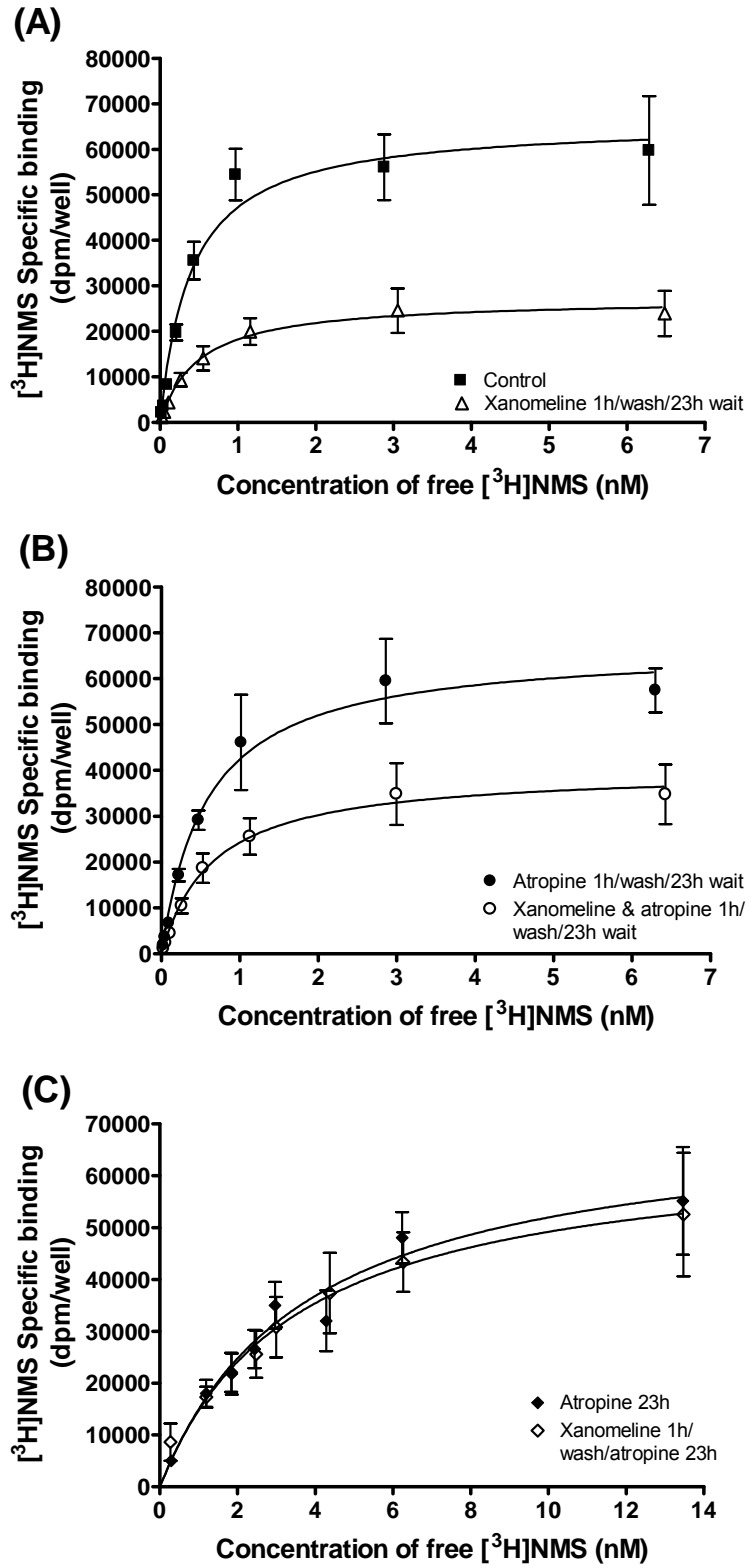
To further explore the importance of receptor activation in the long-term effects of xanomeline at the M3 receptor, receptor function was obliterated using siRNA targeted against the G<sub>q</sub> and G<sub>11</sub> G proteins (Fig. 7). Functional assays and western blots confirmed that receptor response and G<sub>q/11</sub> G protein expression were significantly reduced. When siRNA-transfected cells were subjected to the various xanomeline treatments, no differences in receptor density on the cell-surface were observed. This is in contrast to non-transfected cells in which prolonged xanomeline pretreatments resulted in decreased cell-surface receptor density. These results further support the importance of receptor function in the development of xanomeline's long-term effects. In contrast, treatment with xanomeline for 24 hours followed by washing resulted in a decrease in radioligand affinity at both siRNA-treated and untreated M3 receptors, further supporting the notion that xanomeline binding can initiate changes in radioligand affinity in the absence of function. This further suggests allosteric regulation of the primary binding site by wash-resistant xanomeline binding.

Previous experiments at the M1 receptor have shown that changing the structure of xanomeline, specifically shortening the carbon side chain, results in alterations in its persistent binding and receptor activation (Jakubik et al., 2004). We confirmed that the propyl analog of xanomeline, which has three fewer carbons in the side-chain, exhibits low-potency wash-resistant binding at the M3 receptor in the virtual absence of concomitant receptor activation. Using [<sup>3</sup>H]NMS saturation binding paradigms (Fig. 8), pretreatment with the propyl analog for one hour followed by washing and prolonged incubation resulted in a very small decrease in cell-surface receptor density compared to that elicited by xanomeline, without a change in radioligand affinity. In contrast, 24-hour treatment with the propyl analog prior to washing resulted in an unexpected significant ( $p < 0.05$ ) *increase* in cell-surface receptor density in conjunction with a large reduction in

radioligand affinity. This is in stark contrast to the effects induced by xanomeline treatment for 24 hours prior to washing, whereby a marked decrease in cell-surface receptor density was observed. Taken together, these results suggest that wash-resistant agonist binding in the absence of persistent receptor activation is not the major contributor to the long-term effects of xanomeline binding.

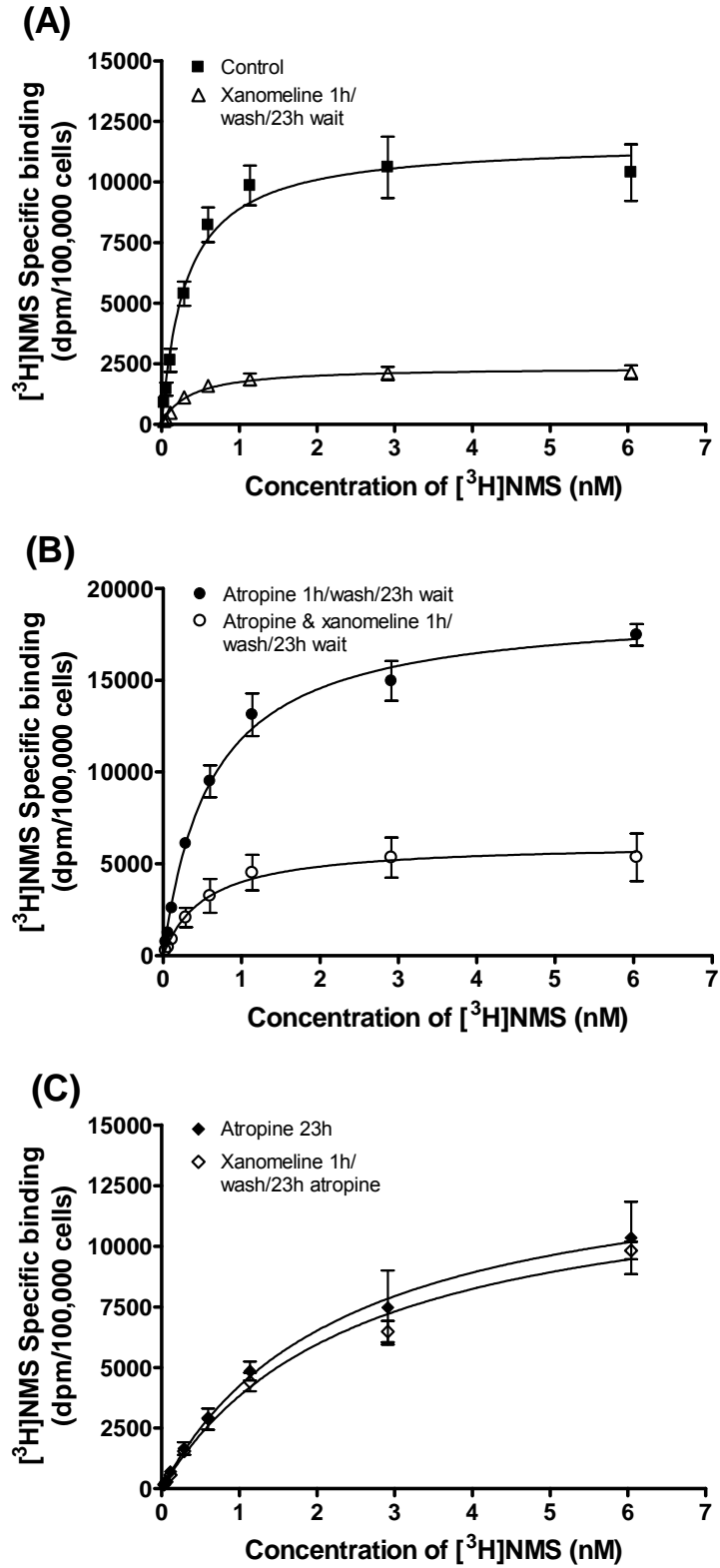
In summary, xanomeline has previously been shown to bind in both a reversible and wash-resistant manner at the M1 and M3 receptors. This binding followed by prolonged exposure results in long-term changes in the binding and functional properties at both receptor subtypes. These results demonstrate that receptor activation is necessary for the majority of the long-term receptor regulation by xanomeline. However, wash-resistant xanomeline binding is able to allosterically modulate the primary binding site, even in the absence of receptor function.

Figure 1



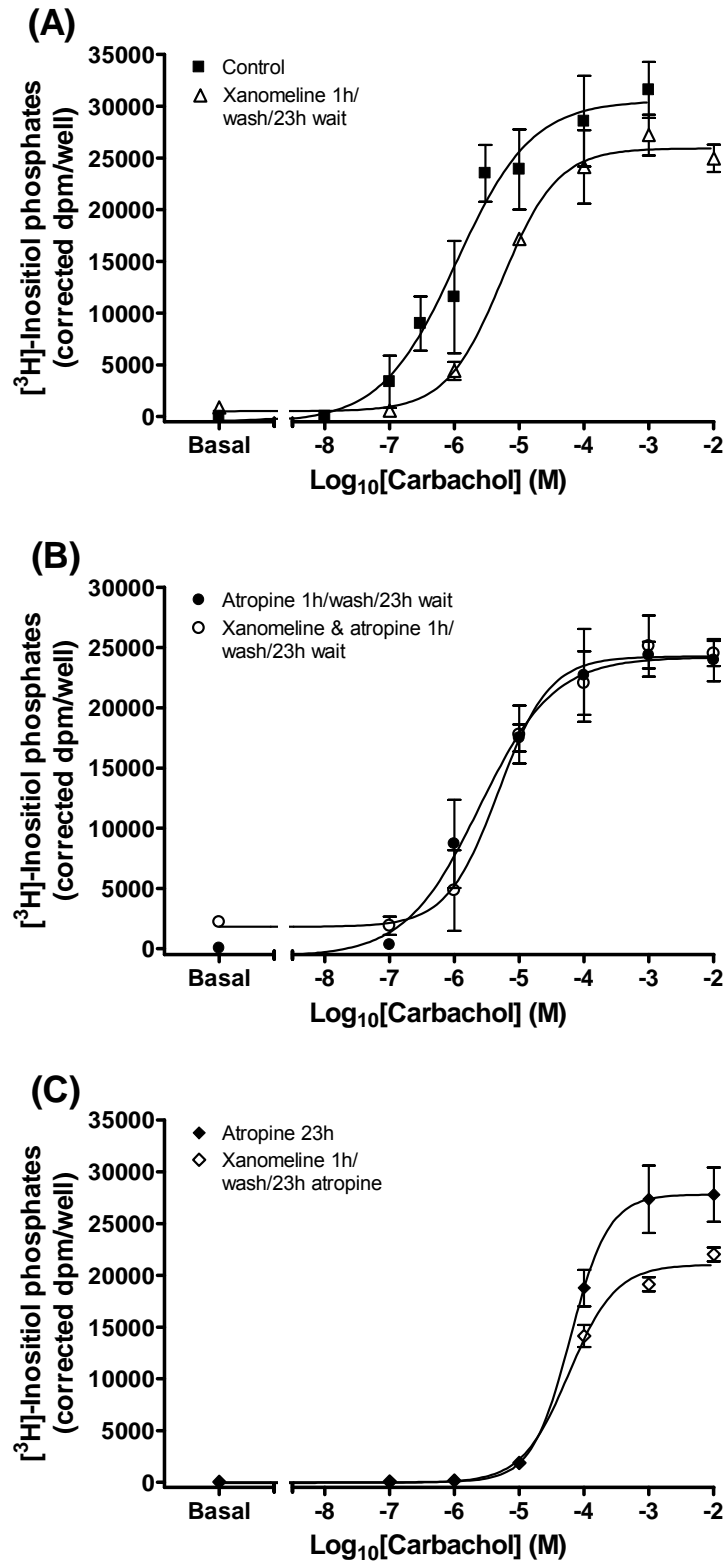
**Figure 1. Effects of atropine on long-term xanomeline-induced inhibition of [<sup>3</sup>H]NMS binding in CHO cells expressing hM3 receptors.** A) Cells were either untreated (closed squares) or treated with 10 μM xanomeline for 1 hour followed by washing and waiting for 23 hours (open triangles). B) Cells were treated with 10 μM atropine for 1 hour followed by washing and waiting 23 hours in control medium (closed circles) or treated simultaneously with atropine and xanomeline for 1 hour followed by washing and waiting 23 hours in control medium (open circles). C) Cells were treated with atropine for 23 hours prior to washing (closed diamonds) or with xanomeline for 1 hour followed by washing and incubation with atropine for 23 hours prior to washing (open diamonds). Saturation binding assays were conducted with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 14 nM) for 1 hour at 37°C. Values represent the means ± S.E. of 4 experiments conducted in duplicate.

Figure 2



**Figure 2. Effects of atropine on long-term xanomeline-induced inhibition of [<sup>3</sup>H]NMS binding in CHO cells expressing hM1 receptors.** A) Cells were either untreated (closed squares) or treated with 3 μM xanomeline for 1 hour followed by washing and waiting for 23 hours (open triangles). B) Cells were treated with 10 μM atropine for 1 hour followed by washing and waiting 23 hours in control medium (closed circles) or treated simultaneously with atropine and xanomeline for 1 hour followed by washing and waiting 23 hours in control medium (open circles). C) Cells were treated with atropine for 23 hours prior to washing (closed diamonds) or with xanomeline for 1 hour followed by washing and incubation with atropine for 23 hours prior to washing (open diamonds). Saturation binding assays were conducted in suspension with increasing concentrations of [<sup>3</sup>H]NMS (0.02 – 4.5 nM) for 1 hour at 37°C. Values represent the means ± S.E. of 3 experiments conducted in triplicate. Data provided by Marianne Grant.

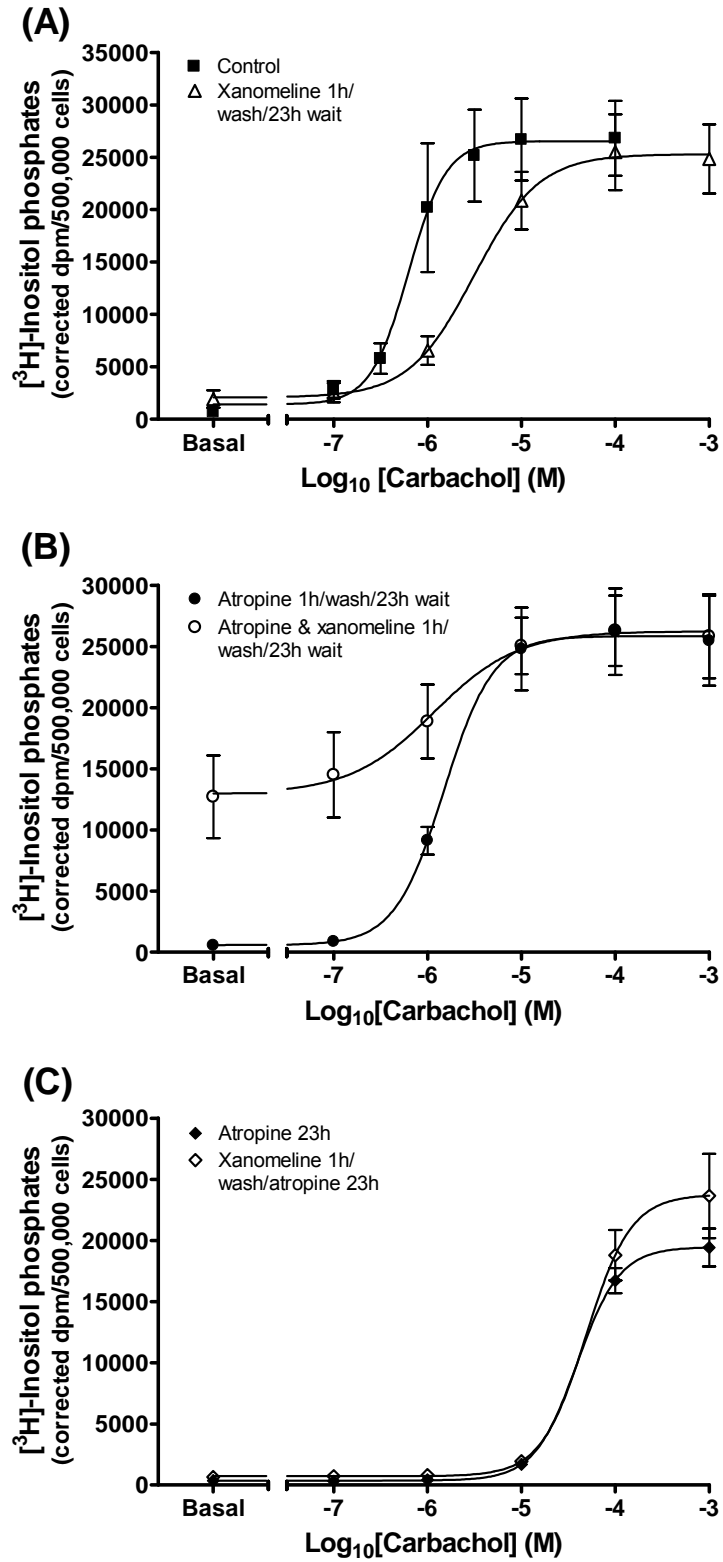
**Figure 3**



**Figure 3. Effects of atropine on long-term modulation of inositol phosphates production by xanomeline in CHO cells expressing hM3 receptors.** A) Cells were either untreated (closed square) or treated with 10  $\mu$ M xanomeline for 1 hour followed by washing and waiting for 23 hours in control medium (open triangles). B) Cells were treated with atropine (10  $\mu$ M) for 1 hour (closed circles) or treated simultaneously with atropine and xanomeline for 1 hour (open circles) followed by washing and waiting 23 hours in control medium. C) Cells were untreated (closed diamonds) or treated with xanomeline for 1 hour (open diamonds) followed by washing and incubation with atropine for 23 hours prior to washing. In all cases, the production of inositol phosphates was determined after the various pretreatments in the presence of increasing concentrations of carbachol (10 nM – 1 mM) for 1 hour at 37°C in the presence of 10 mM LiCl. Values represent the means  $\pm$  S.E. of 3 experiments conducted in triplicate.

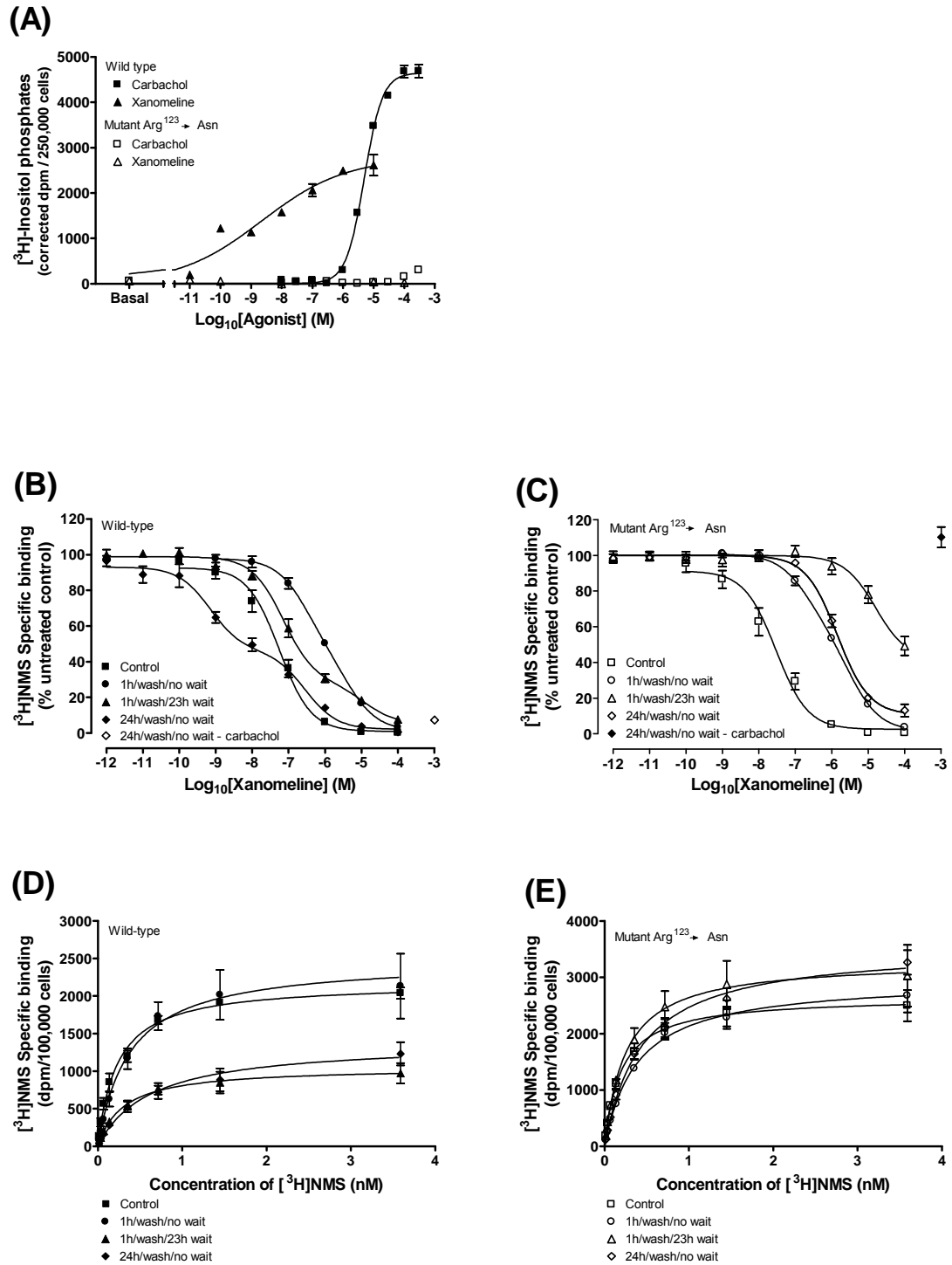


# Figure 4



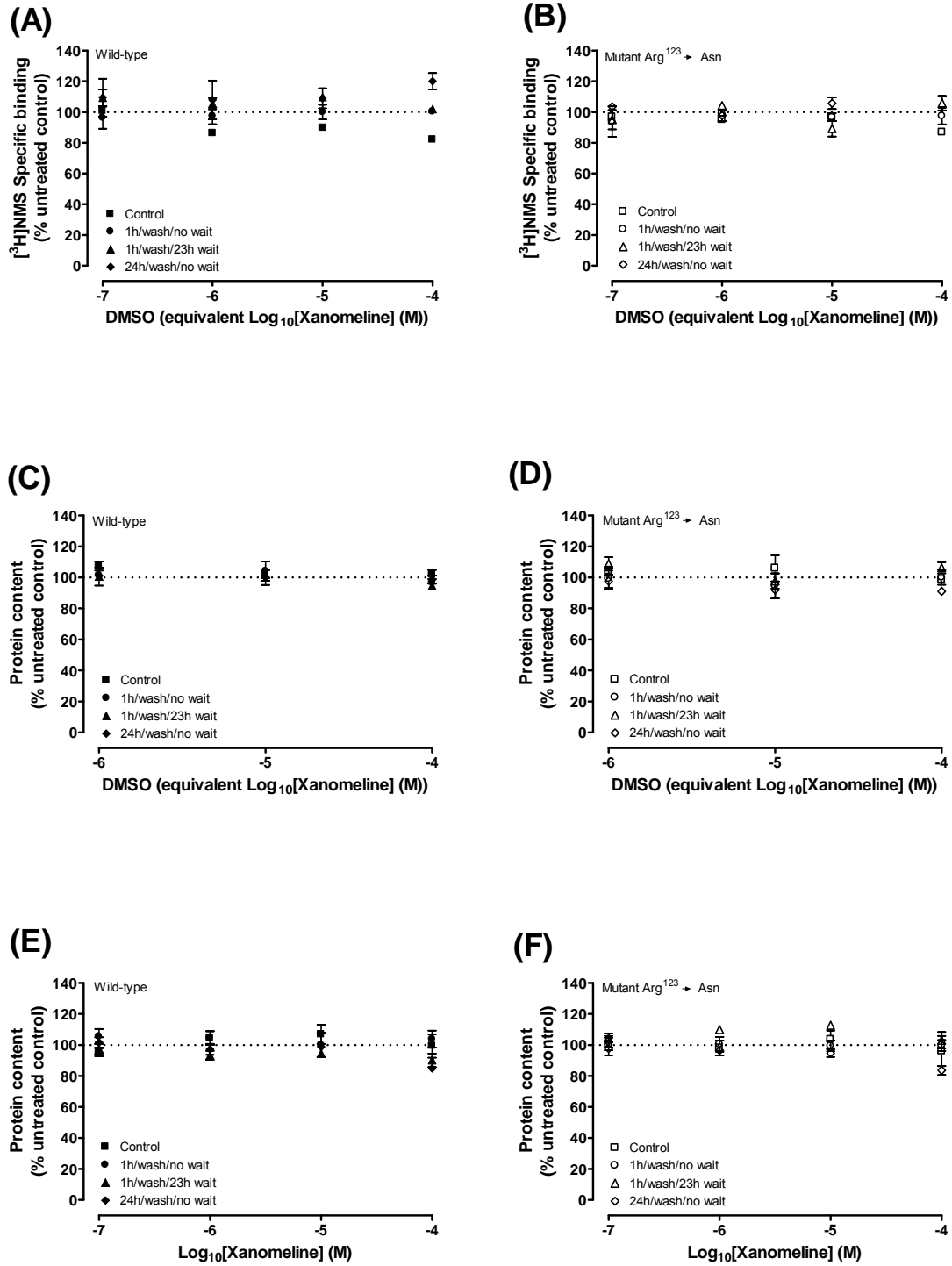
**Figure 4. Effects of atropine on long-term modulation of inositol phosphates production by xanomeline in CHO cells expressing hM1 receptors.** A) Cells were either untreated (closed square) or treated with 300 nM xanomeline for 1 hour followed by washing and waiting for 23 hours in control medium (open triangles). B) Cells were treated with atropine (10  $\mu$ M) for 1 hour (closed circles) or treated simultaneously with atropine and xanomeline for 1 hour (open circles) followed by washing and waiting 23 hours in control medium. C) Cells were untreated (closed diamonds) or treated with xanomeline for 1 hour (open diamonds) followed by washing and incubation with atropine for 23 hours prior to washing. In all cases, the production of inositol phosphates was determined after the various pretreatments in the presence of increasing concentrations of carbachol (10 nM – 1 mM) for 1 hour at 37°C in the presence of 10 mM LiCl. Values represent the means  $\pm$  S.E. of 3-4 experiments conducted in triplicate. Data provided by Kayla De Lorme and Marianne Grant.

# Figure 5



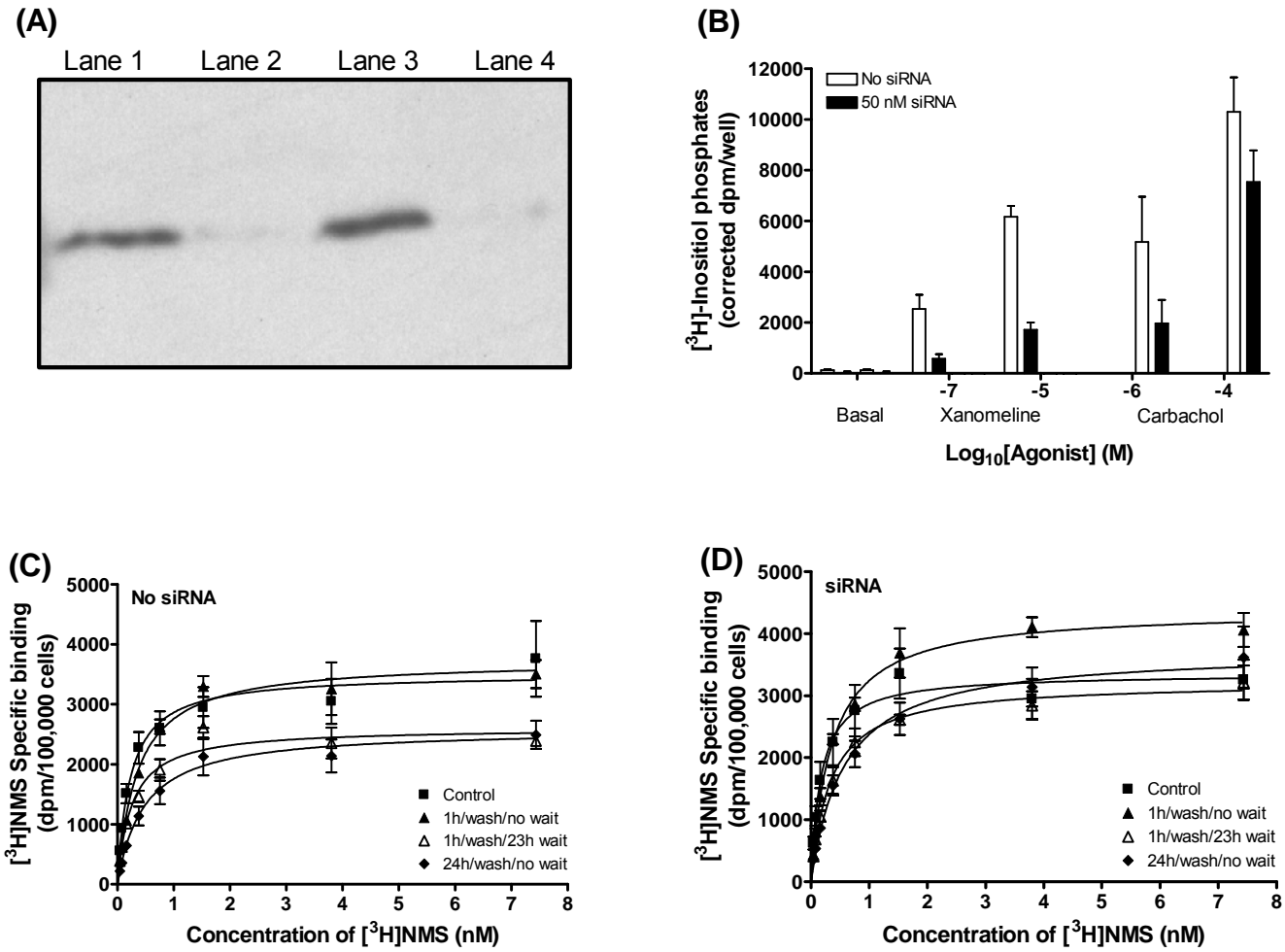
**Figure 5. Effects of using a non-functional receptor mutant on function, inhibition of [<sup>3</sup>H]NMS and saturation binding in CHO cells expressing rM1 receptors.** A) Cells expressing wild-type (closed symbols) and mutant rM1 (open symbols) receptors were stimulated with increasing concentrations of carbachol (10 nM – 100 μM) (squares) or xanomeline (10 pM – 10 μM) (triangles) for 1 hour at 37°C in the presence of 10 mM lithium chloride and inositol phosphate production was determined. B, D) Cells expressing wild-type rM1 receptors were treated with xanomeline for 1 hour followed by washing and used immediately (circles) or subsequent to 23 hour incubation in control media (triangles). Additional cells were treated with xanomeline for 24 hours prior to washing (diamonds). B) Untreated cells were incubated concurrently with increasing concentrations of xanomeline and 0.2 nM [<sup>3</sup>H]NMS or xanomeline pretreated cells were incubated with 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37 °C. D) Untreated or xanomeline pretreated cells were incubated with increasing concentration of [<sup>3</sup>H]NMS (0.02 -3.5 nM) for 1 hour at 37°C. C, E) Cells expressing the rM1<sup>123</sup> mutant receptor were treated the same as described for (B) and (D). For B and C individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 13100 and 11500 dpm/well for wild-type and mutant rM1 receptors, respectively. Values represent the means ± S.E. of 2-7 experiments conducted in triplicate. Data in (A) provided by Marianne Grant.

Figure 6



**Figure 6. Control experiments to determine [<sup>3</sup>H]NMS binding and protein content following xanomeline or DMSO pretreatments in CHO cells expressing rM1 receptors.** Cells expressing wild-type (closed symbols) or rM1<sup>123</sup> mutant receptors (open symbols) were treated with increasing concentrations of xanomeline or the corresponding concentrations of DMSO for 1 hour followed by washing and used immediately (circles) or subsequent to 23 hour incubation (triangles). Additional cells were treated with xanomeline or the corresponding concentration of DMSO for 24 hours prior to washing (diamonds). Control cells were untreated (squares). A, B) Following pretreatment with DMSO cells were incubated with 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. C, D) Following pretreatment with DMSO protein content was determined using the method of Bradford (1976). E, F) Following pretreatment with xanomeline protein content was determined using the method of Bradford (Bradford, 1976). For A and B individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 9300 and 7500 dpm/well for wild-type and mutant rM1 receptors, respectively. For C and D individual experiments were normalized to control protein in the absence of DMSO treatment which had a mean of 227 and 175 µg/well for wild-type and mutant rM1 receptors, respectively. For E and F individual experiments were normalized to control protein in the absence of xanomeline treatment which had a mean of 223 and 185 µg/well for wild-type and mutant rM1 receptors, respectively. Values represent the means ± S.E. of 2-4 experiments conducted in duplicate or triplicate

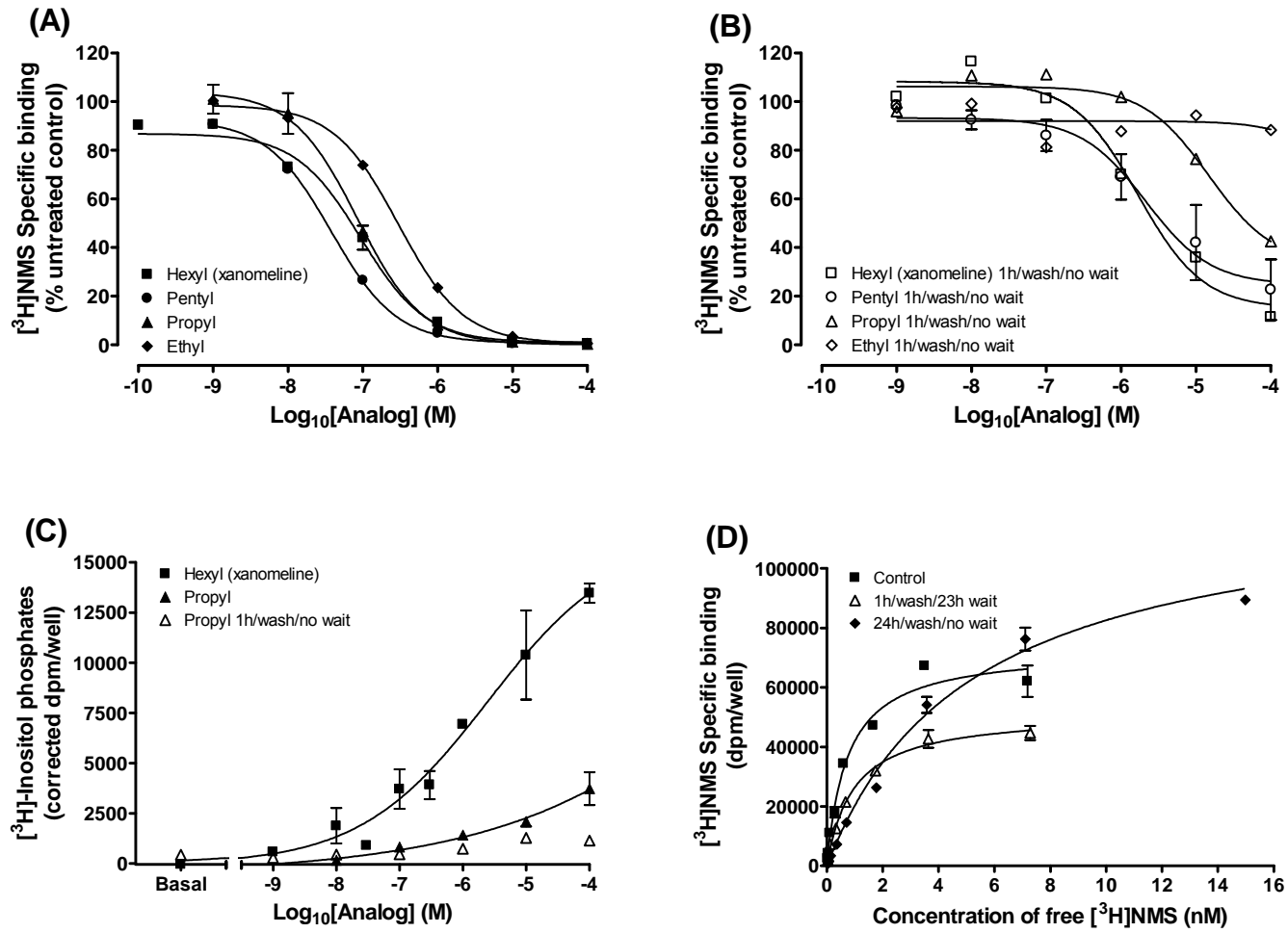
Figure 7



**Figure 7. Effects of blocking receptor function using siRNA against G<sub>q</sub> and G<sub>11</sub> G proteins on long-term xanomeline-induced inhibition of [<sup>3</sup>H]NMS binding in CHO cells expressing the hM3 receptor.** A) Cells were left untreated (lane 1 & 3) or treated with 50 nM siRNA for G<sub>q</sub> and G<sub>11</sub> G proteins (lane 2 & 4). The level of G<sub>q/11</sub> G protein expression was determined by western blots. B) Cells were untreated (open bars) or treated with 50 nM siRNA for G<sub>q</sub> and G<sub>11</sub> G proteins (solid bars). Inositol phosphate production was determined following stimulation with two concentrations of carbachol (1 μM or 100 μM) or xanomeline (100 nM or 10 μM). C) Cells were treated with xanomeline (10 μM) for 1 hour followed by washing and used immediately (closed triangles) or after 23 hour incubation in control medium (open triangles). An additional group of cells was treated with xanomeline for 24 hours prior to washing (closed diamonds). Subsequently, untreated cells (closed squares) or pretreated cells were incubated in suspension with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 7.5 nM) for 1 hour at 37°C. D) Cells were treated with 50 nM siRNA for G<sub>q</sub> and G<sub>11</sub> G proteins to inhibit receptor function then subjected to various xanomeline treatment protocols as in (C). Values represent the means ± S.E. of 3 experiments conducted in triplicate.



Figure 8



**Figure 8. Effects of wash-resistant binding in the absence of receptor activation on the ability of the propyl analog of xanomeline to initiate long-term effects on saturation binding parameters in CHO cells expressing hM3 receptors.**

The xanomeline analogs that were studied are hexyl (xanomeline; squares), pentyl (circles), propyl (triangles) and ethyl (diamonds). A) Cells were exposed to xanomeline analogs concurrently with 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37°C (closed symbols). B) Cells were pretreated with increasing concentrations of xanomeline analogs for 1 hour followed by washing (open symbols). Subsequently, cells were exposed to 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. C) Cells were stimulated with increasing concentrations of xanomeline (closed squares) or the propyl analog (closed triangles) for 1 hour in the presence of 10 mM LiCl and inositol phosphate production was determined. Additional cells were pretreated with the propyl analog for 1 hour followed by washing (open triangles). The production of inositol phosphates was determined in the absence of further agonist stimulation in the presence of 10 mM LiCl. D) Cells were pretreated with 100 μM of the propyl xanomeline analog for 1 hour followed by washing and incubation in the absence of free drug for 23 hours (open triangles). An additional group of cells was treated for 24 hours with the propyl analog prior to washing (closed diamonds). Subsequently, untreated cells (closed square) or pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.02 - 15 nM) for 1 hour at 37°C. For A and B individual experiments were normalized to control binding in the absence of treatment with xanomeline analogs which had a mean of 14400 dpm/well. Values represent the means ± S.E. of 3 experiments conducted in duplicate.

**Table 1. Effects of xanomeline pretreatment and washout in the absence or presence of atropine on [<sup>3</sup>H]NMS saturation binding and functional parameters of carbachol stimulation in CHO cells expressing hM1 or hM3 receptors.** Cells were pretreated for 1 hour with 3 μM (M1, binding), 300 nM (M1, function) or 10 μM (M3) xanomeline in the absence or presence of 10 μM atropine followed by washing and waiting for 23 hours in the absence or presence of atropine. In the binding assay, untreated or pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS for 1 hour at 37°C. In the functional assay, untreated or pretreated cells were stimulated with increasing concentrations of carbachol for 1 hour at 37°C in the presence of 10 mM lithium chloride. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

	<u>Saturation Binding Parameters</u>		<u>Functional Parameters</u>	
	$K_d^a$ (nM)	$B_{max}^b$	$pEC_{50}^c$	$E_{max}^d$
<b>M1 receptor – pretreatment groups</b>				
Control (untreated) <sup>e</sup>	0.31 ± 0.02	11600 ± 1200	6.1 ± 0.09	24700 ± 2200
1 h xanomeline/washout/23 h wait	0.36 ± 0.02	2400 ± 300 <sup>s</sup>	5.4 ± 0.12 <sup>s</sup>	23500 ± 2100
1 h atropine/washout/23 h wait	0.52 ± 0.09	19000 ± 800	5.9 ± 0.09	26600 ± 3400
1 h xanomeline & atropine/washout/23 h wait	0.55 ± 0.02	6100 ± 1300 <sup>#</sup>	5.9 ± 0.10	25800 ± 3600
23 h atropine/washout/no wait	2.38 ± 0.31	14200 ± 2700	4.4 ± 0.07	19500 ± 1600
1h xanomeline/washout/23 h atropine/washout/no wait	2.64 ± 0.49	13500 ± 800	4.3 ± 0.04	23800 ± 3500
<b>M3 receptor – pretreatment groups</b>				
Control (untreated) <sup>e</sup>	0.38 ± 0.05	66000 ± 11000	6.0 ± 0.18	29600 ± 2700
1 h xanomeline/washout/23 h wait	0.47 ± 0.03*	27000 ± 5500*	5.3 ± 0.07*	26100 ± 2300
1 h atropine/washout/23 h wait	0.58 ± 0.05	67000 ± 7400	5.6 ± 0.17	24200 ± 2000
1 h xanomeline & atropine/washout/23 h wait	0.66 ± 0.07	40000 ± 7400 <sup>†</sup>	5.3 ± 0.16	24600 ± 1600
23 h atropine/washout/no wait	3.20 ± 0.39	68000 ± 11800	4.2 ± 0.01	28200 ± 3200
1h xanomeline/washout/23 h atropine/washout/no wait	3.25 ± 0.71	67000 ± 17000	4.2 ± 0.09	21100 ± 830

Table 1 cont.

<sup>a</sup> Equilibrium dissociation constant for [<sup>3</sup>H]NMS binding.

<sup>b</sup> Maximal cell-surface receptor density (dpm/100,000 cells (M1) or dpm/well (M3)).

<sup>c</sup> Negative logarithm of midpoint (potency) parameter.

<sup>d</sup> Maximal response. Values are expressed as dpm/500,000 cells (M1) or dpm/well (M3) after correcting for standard recovery.

<sup>e</sup> Control experiments were conducted on untreated cells in the binding assay and untreated cells were stimulated with carbachol in the functional assay.

<sup>§</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in  $K_d$ ,  $B_{max}$ ,  $pEC_{50}$  or  $E_{max}$  where indicated between the 1h/wash/23h wait xanomeline pretreated group compared to control at the M1 receptor.

<sup>#</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in  $K_d$ ,  $B_{max}$ ,  $pEC_{50}$  or  $E_{max}$  where indicated between the 1h xanomeline and atropine/wash/23h wait pretreated group compared to 1h atropine/wash/23h wait treatment at the M1 receptor.

\*Paired t-test detected a significant difference ( $p < 0.05$ ) in  $K_d$ ,  $B_{max}$ ,  $pEC_{50}$  or  $pEC_{50}$  where indicated between the 1h/wash/23h wait xanomeline pretreated group compared to control at the M3 receptor.

<sup>†</sup> Paired t-test detected a significant difference ( $p < 0.05$ ) in  $K_d$ ,  $B_{max}$ ,  $pEC_{50}$  or  $E_{max}$  where indicated between the 1h xanomeline and atropine/wash/23h wait pretreated group compared to 1h atropine/wash/23h wait treatment at the M3 receptor.

No significant difference was detected between the 1h xanomeline/wash/23h atropine/wash/no wait group compared to 23h atropine/wash/no wait treatment at either the M1 or M3 receptor.

**Table 2. Effects of xanomeline pretreatment on inhibition of NMS binding, saturation binding and functional parameters in wild-type and mutant rM1 receptors expressed in CHO cells.** Cells were pretreated with increasing concentrations (1 pM - 100µM) or 3 µM xanomeline for one hour followed by washing and used immediately or after 23 hour incubation in ligand free media. Additional cells were treated with increasing concentrations (1 pM – 100 µM) or 3 µM xanomeline for 24 hour prior to washing. In the functional assay, untreated cells were exposed to increasing concentrations of carbachol or xanomeline in the presence of 10 mM lithium chloride for one hour at 37°C. In the [<sup>3</sup>H]NMS inhibition binding assay, untreated cells were incubated simultaneously with xanomeline and 0.2 nM [<sup>3</sup>H]NMS or pretreated cells were incubated with 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. In the saturation binding assays, untreated or pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.02 – 3.6 nM) for 1 hour at 37°C. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

	rM1 wild-type							
	<u>Functional parameters</u>		<u>Inhibition of [<sup>3</sup>H]NMS binding parameters</u>				<u>Saturation parameters</u>	
	pEC <sub>50</sub> <sup>a</sup>	E <sub>max</sub> <sup>b</sup>	pIC <sub>50</sub> <sup>c</sup>	pIC <sub>50</sub> – H <sup>d</sup>	pIC <sub>50</sub> – L <sup>e</sup>	I <sub>max</sub> <sup>f</sup>	K <sub>d</sub> <sup>g</sup> (nM)	B <sub>max</sub> <sup>h</sup>
<b>Carbachol</b>								
Control (untreated)	5.3	4700						
<b>Xanomeline</b>								
Control (untreated)	8.7	2800	7.3 ± 0.14			99 ± 0.6	0.27 ± 0.08	2200 ± 90
1 h/washout/no wait			6.0 ± 0.06 <sup>s</sup>			95 ± 0.5	0.40 ± 0.04	2600 ± 400
1 h/washout/23 h wait				7.1 ± 0.16* (74 ± 1.1%)	4.6 ± 0.23*	99 ± 2.3	0.30 ± 0.04	1100 ± 150 <sup>#</sup>
24 h/washout/no wait				9.4 ± 0.20* (47 ± 7.7%)	6.6 ± 0.12	97 ± 0.5	0.60 ± 0.12 <sup>#</sup>	1300 ± 150

Table 2 cont.

	rM1 <sup>123</sup> Mutant							
	<u>Functional parameters</u>		<u>Inhibition of [<sup>3</sup>H]NMS binding parameters</u>				<u>Saturation parameters</u>	
	pEC <sub>50</sub>	E <sub>max</sub>	pIC <sub>50</sub>	pIC <sub>50</sub> – H	pIC <sub>50</sub> – L	I <sub>max</sub>	K <sub>d</sub> (nM)	B <sub>max</sub> <sup>h</sup>
<b>Carbachol</b>								
Control (untreated)	ND	ND						
<b>Xanomeline</b>								
Control (untreated)	ND	ND	7.6 ± 0.14			99 ± 0.5	0.19 ± 0.02	2700 ± 300
1 h/washout/no wait			6.0 ± 0.02 <sup>§</sup>			96 ± 0.7	0.42 ± 0.10	3000 ± 380
1 h/washout/23 h wait			4.9 ± 0.23 <sup>*</sup>			60 ± 7.4 <sup>‡</sup>	0.26 ± 0.07	3400 ± 600
24 h/washout/no wait			5.8 ± 0.08			90 ± 2.9	0.49 ± 0.11	3600 ± 320

<sup>a</sup> Negative logarithm of midpoint (potency) parameter.

<sup>b</sup> Maximal response. Values are expressed as dpm/250,000 cells after correcting for standard recovery.

<sup>c</sup> Negative logarithm of the IC<sub>50</sub> for binding to a single potency site.

<sup>d</sup> Negative logarithm of the IC<sub>50</sub> for the high-potency site. Percent of binding sites shown in parentheses.

<sup>e</sup> Negative logarithm of the IC<sub>50</sub> for the low-potency site.

<sup>f</sup> Maximal percent inhibition of [<sup>3</sup>H]NMS binding.

<sup>g</sup> Equilibrium dissociation constant for [<sup>3</sup>H]NMS binding.

<sup>h</sup> Maximal cell-surface receptor density (dpm/100,000 cells).

Statistical analysis was not performed for functional parameters (pEC<sub>50</sub> or E<sub>max</sub>) because the data presented are from a single experiment conducted in duplicate.

<sup>§</sup> Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> between the xanomeline 1h/wash/no wait group compared to their respective controls at the wild-type or mutant rM1 receptor.

<sup>\*</sup>ANOVA followed by Dunnett's post-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> between the 1h/wash/23h wait or 24h/wash/no wait groups compared their respective 1h/wash/no wait group at the wild-type or mutant rM1 receptor.

<sup>#</sup>ANOVA followed by Dunnett's post-test detected a significant difference ( $p < 0.05$ ) in K<sub>d</sub> or B<sub>max</sub> where indicated between the xanomeline pretreated groups compared to their respective controls at the wild-type or mutant rM1 receptor.

<sup>‡</sup> Significantly different ( $p < 0.5$ ) from 100 percent as determined by ANOVA followed by Dunnett's post-test comparison.

**Table 3. Effects of blocking receptor function using siRNA against G<sub>q</sub> and G<sub>11</sub> G proteins on [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing hM3 receptors assayed in suspension.** Cells were untreated or treated with 50 nM siRNA for G<sub>q</sub> and G<sub>11</sub> G proteins. Subsequently, cells were treated with 10 μM xanomeline for 1 hour followed by washing and used immediately or after 23 hour incubation in control media. An additional group of cells was treated with xanomeline for 24 hours prior to washing. Cells were then incubated in suspension with increasing concentrations of [<sup>3</sup>H]NMS (0.04 – 7.4 nM) at 37°C for 1 hour. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

Xanomeline pretreatment group	Control (no siRNA)		siRNA	
	K <sub>d</sub> <sup>a</sup> (nM)	B <sub>max</sub> <sup>b</sup>	K <sub>d</sub> (nM)	B <sub>max</sub>
Control (untreated)	0.22 ± 0.05	3500 ± 500	0.22 ± 0.06	3400 ± 300
1 h/washout/no wait	0.33 ± 0.04	3800 ± 400	0.40 ± 0.07	4400 ± 300
1 h/washout/23 h wait	0.24 ± 0.02	2600 ± 100	0.36 ± 0.01	3200 ± 200
24 h/washout/no wait	0.48 ± 0.12	2600 ± 300	0.63 ± 0.13*	3700 ± 500

<sup>a</sup> Equilibrium dissociation constant of [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell-surface receptor density (dpm/100,000 cells).

\*ANOVA followed by Dunnett's post-test detected a significant difference ( $p < 0.05$ ) in K<sub>d</sub> or B<sub>max</sub> where indicated between the siRNA-pretreated groups compared with siRNA treated control.

No significant difference was detected between no siRNA-pretreated groups compared to no siRNA treated control.

**Table 4. Effects of xanomeline analogs on inhibition of [<sup>3</sup>H]NMS binding and functional parameters in CHO cells expressing hM3 receptors.** Cells were treated with increasing concentration of the xanomeline analogs (1 nM – 100 μM) for 1 hour followed by washing. In the binding assay, untreated cells were incubated concurrently with 0.2 nM [<sup>3</sup>H]NMS and xanomeline analogs or pretreated cells were incubated with 0.2 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. In the functional assay, untreated cells were stimulated with increasing concentration of the analogs and pretreated cells were not further stimulated in the presence of 10 mM lithium chloride for 1 hour at 37°C.

	<u>Hexyl (xanomeline)</u>				<u>Pentyl</u>		<u>Propyl</u>				<u>Ethyl</u>	
	Binding parameters		Functional parameters		Binding parameters		Binding parameters		Functional parameters		Binding parameters	
	pIC <sub>50</sub> <sup>a</sup>	I <sub>max</sub> <sup>b</sup>	pEC <sub>50</sub> <sup>c</sup>	E <sub>max</sub> <sup>d</sup>	pIC <sub>50</sub>	I <sub>max</sub>	pIC <sub>50</sub>	I <sub>max</sub>	pEC <sub>50</sub>	E <sub>max</sub>	pIC <sub>50</sub>	I <sub>max</sub>
Control (untreated)	7.0 ± 0.02	99 ± 0.2	6.1 ± 0.24	14500	7.4	99	7.1 ± 0.01	100 ± 0.1	3.4	6500	6.6	99
1h /wash/no wait	5.8	85			5.9	87	5.1 ± 0.26	66 ± 0.8	ND	ND	ND	ND

<sup>a</sup> Negative logarithm of the IC<sub>50</sub> for binding to a single potency site.

<sup>b</sup> Maximal percent inhibition of [<sup>3</sup>H]NMS binding.

<sup>c</sup> Negative logarithm of midpoint (potency) parameter.

<sup>d</sup> Maximal response. Values are expressed as dpm/well after correcting for standard recovery.

ND – parameter values were not determined

Statistical analyses were not performed because data are the results of 1-2 experiments conducted in duplicate or triplicate.



**Table 5. Effects of treatment with the propyl xanomeline analog on [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing hM3 receptors.** Cells were treated with 100 μM of the propyl analog for 1 hour followed by washing and used after 23 hour incubation in control media or treated with the propyl analog for 24 hours prior to washing. Untreated or pretreated cells were then incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.03 – 7.2 nM) for 1 hour at 37°C. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

<b>Pretreatment group</b>	<b>K<sub>d</sub><sup>a</sup> (nM)</b>	<b>B<sub>max</sub><sup>b</sup></b>
Control (untreated)	0.74 ± 0.10	73400 ± 4900
1 h/washout/23h wait	0.93 ± 0.06	51500 ± 3100
24 h/washout/no wait	5.73 ± 0.74*	133100 ± 10500*

<sup>a</sup> Equilibrium dissociation constant of [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell-surface receptor density (dpm/well).

\*ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in K<sub>d</sub> and B<sub>max</sub> where indicated between the propyl analog-pretreated groups compared with control.

**CHAPTER 6: Mechanisms of the long-term effects of xanomeline wash-resistant binding.**

## **Introduction**

Acute wash-resistant binding of xanomeline to muscarinic receptors leads to delayed changes in receptor binding and functional properties. Previous research has shown that receptor regulation following prolonged exposure to conventional agonists can occur through various mechanisms such as internalization or down-regulation of the receptor or uncoupling of the receptor from the G protein (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990). In addition these changes have been shown to occur in a time-dependent manner. It has been shown that brief treatment of cells that express muscarinic receptors with carbachol results in a decrease in binding of hydrophilic ligands (e.g., [<sup>3</sup>H]NMS), but not lipophilic ones (e.g. [<sup>3</sup>H]QNB) (Maloteaux et al., 1983). These ligands are capable of binding to only cell surface receptors and both cell surface and internalized receptors, respectively. Thus, brief treatment with carbachol induces receptors to internalize without being down regulated. In contrast long-term carbachol treatment causes receptor down regulation.

Many factors are involved in agonist-induced receptor regulation, including the preferential signal pathway to which the receptor is coupled, agonist efficacy and temperature. Both the M1 and M3 receptors are coupled to the G<sub>q/11</sub> G proteins which lead to stimulation of phospholipase C (Volpicelli & Levey, 2004). In contrast, the M2 and M4 receptors are coupled to the G<sub>i/o</sub> G proteins that lead to the inhibition of adenylate cyclase (Eglen, 2005). This may influence their susceptibility to regulation by agonists. Receptor regulation is also proportional to the relative efficacy of agonists at a given receptor. However, it should be noted that receptor overexpression in transfected cells could endow a partial agonist with full efficacy (Kenakin, 1997; Wood et al., 1999). It has also been shown that decreasing the temperature inhibits receptor function and internalization (El-Fakahany & Richelson, 1980; Maloteaux & Hermans, 1994).

Following prolonged exposure, xanomeline has been shown to result in a decrease in receptor expression and antagonism of the functional response to subsequent addition of agonists. The changes in the long-term responses appear to be similar at the M1 and M3 receptors despite the fact that xanomeline is a full agonist at the M1 receptor, but a partial agonist at the M3 receptor. Thus, experiments were designed to: 1) differentiate the effects of xanomeline pretreatment on receptor internalization and down regulation; 2) test the effects of changing receptor expression level on xanomeline effects; 3) further assess the role of receptor signaling in the ability of brief pretreatment with xanomeline to induce delayed receptor regulation.

## **Results**

### **Comparison of the effects of brief and long-term exposure of cells to xanomeline on [<sup>3</sup>H]QNB and [<sup>3</sup>H]NMS binding at the M3 and M1 receptors**

Prolonged agonist exposure can lead to receptor internalization, down-regulation or uncoupling of the receptor from its G protein (Dohlman et al., 1991; El-Fakahany & Cioffi, 1990). The current set of experiments was conducted to differentiate the long-term effects of xanomeline on internalization and down-regulation of the M3 and M1 receptors. These receptors both couple to G<sub>q/11</sub> G proteins (Volpicelli & Levey, 2004), resulting in the possibility that the mechanism of regulation may be similar between these two subtypes. QNB is a lipophilic muscarinic receptor antagonist that is able to bind to both cell-surface and internalized (but intact) receptors (Ehlert et al., 1981; Vickroy & Malphurs, 1994). In contrast, NMS (a permanently-charged quaternary amine) is only able to bind to receptors expressed on the cell surface (Feigenbaum & El-Fakahany, 1985). We compared effects of the various xanomeline treatment paradigms on the specific binding of both radioligands. Receptor-saturating concentrations of

[<sup>3</sup>H]QNB and [<sup>3</sup>H]NMS were used to confine effects of xanomeline to alterations in the number of receptors without interference from changes in radioligand affinity.

Cells were exposed to increasing concentrations of xanomeline for the various treatment conditions and the binding of either 1.4 nM [<sup>3</sup>H]QNB or 2.8 nM [<sup>3</sup>H]NMS was determined. These concentrations were chosen because they result in similar receptor occupancy by either ligand (approximately 93% of total receptors). In untreated cells expressing the M3 receptor, inclusion of xanomeline in the binding assay resulted in complete concentration-dependent inhibition of binding of both radioligands with similar potency (Fig. 1; Table 1). Cells pretreated with xanomeline for one minute (Fig. 1 a, b) or one hour (Fig. 1 c, d) followed by washing and immediate use in the binding assay showed wash-resistant xanomeline binding that exhibited a significantly ( $p < 0.05$ ) lower potency against either radioligand compared to when xanomeline was included in the binding assay (Table 1). Additionally, inhibition of [<sup>3</sup>H]NMS binding was incomplete following this treatment. When cells were pretreated with xanomeline for one minute or one hour followed by washing and waiting 24 (Fig. 1 a, b) or 23 (Fig. 1 c, d) hours, respectively, there was a significant ( $p < 0.05$ ) increase in the apparent potency of xanomeline as compared to its acute wash-resistant effects. Surprisingly, this increase was greater for NMS compared to QNB for one hour xanomeline pretreatment. Noteworthy, inhibition of binding under these treatment conditions was incomplete for both radioligands resulting in a maximal inhibition of approximately 35-45% (Table 1).

Additional groups of cells were incubated with increasing concentrations of xanomeline for 24 hours followed by washing and immediate use in radioligand binding assays. This condition resulted in biphasic xanomeline inhibition curves of either radioligand, with the high-potency site resulting in approximately 40 percent of the binding sites for both radioligands (Fig. 1). The apparent high-potency site under these

conditions demonstrated a markedly lower  $IC_{50}$  than that observed in the case of cells that were briefly treated with xanomeline, washed and used immediately or further incubated in control media for a day (Table 1). Once again, the inhibition of binding was incomplete for both radioligands under these conditions (Table 1).

When parallel experiments were conducted in cells expressing the M1 receptor similar results were obtained for the acute xanomeline treatments. When cells were exposed to increasing concentrations of xanomeline in the concurrent presence of [ $^3$ H]QNB or [ $^3$ H]NMS there was a concentration dependent decrease in binding that resulted in complete inhibition of either radioligand binding and similar potencies of xanomeline (Fig. 2). When cells were exposed to xanomeline for one minute (Fig. 2 a, b) or one hour (Fig. 2 c, d) followed by washing and immediate use there was a concentration-dependent decrease in radioligand binding, however, inhibition was incomplete in all cases (Table 1). It should be noted that the inhibition of [ $^3$ H]QNB binding following one minute pretreatment and washing with xanomeline could not be determined because the binding curve was still decreasing at the highest concentration tested. However, due to technical limitations with the vehicle xanomeline was dissolved in (DMSO) higher concentrations could not be used. Additionally, all acute treatment groups resulted in a significantly ( $p < 0.05$ ) lower potency compared to when xanomeline was continuously present in the binding assay (Table 1). When cells were treated with xanomeline for one minute or one hour followed by washing and waiting 24 (Fig. 2 a, b) or 23 (Fig. 2 c, d) hours, respectively, there was once again a concentration-dependent decrease in radioligand binding that was complete for the one-minute pretreatments, but incomplete for the one-hour pretreatments (Table 1). In all cases there was a significant ( $p < 0.05$ ) increase in the potency compared to the respective acute wash-resistant effects.

In contrast to the results obtained at the M3 receptor, when cells expressing the M1 receptor were exposed to xanomeline for 24 hours prior to washing there was a monophasic decrease in radioligand binding, with similar potencies for both radioligands (Fig. 2). In both cases, the potencies were significantly ( $p < 0.05$ ) higher compared to cells that were briefly treated with xanomeline. Again, the inhibition of radioligand binding was incomplete for both radioligands (Table 1).

### **Effects of wash-resistant binding in the absence of receptor activation on the ability of xanomeline to mediate long-term effects on saturation binding at the M3 receptor**

Previous literature has shown that wash-resistant inhibition of [<sup>3</sup>H]NMS binding by xanomeline occurs with markedly lower potency at 4°C than at 37°C at the M1 receptor (Jakubik et al., 2004). Furthermore, functional assays using cells expressing the M3 receptor demonstrated lack of activation of the receptor by 10 or 100 μM xanomeline at 4°C when it was present for either one or two hours (Fig. 3 a). Therefore, experiments were conducted at 4°C to further demonstrate the need for receptor activation to obtain the long-term effects of xanomeline at the M3 receptor. Increasing concentrations of [<sup>3</sup>H]NMS were used to determine cell-surface receptor density and radioligand affinity. Exposure of cells to xanomeline for one hour at 37°C followed by washing and 23-hour incubation at 4°C in the absence of free xanomeline did not result in changes in maximal binding or radioligand affinity compared to the 4°C control (Fig. 3 b). However, when cells were exposed to 10 or 100 μM xanomeline at 4°C for 24 hours prior to washing there was a significant ( $p < 0.05$ ) concentration-dependent decrease in [<sup>3</sup>H]NMS radioligand affinity in the absence of a change in cell-surface receptor density compared to the 4°C control (Fig. 3 b). Results are summarized in Table 2.

## **Time course of the switch of xanomeline pharmacological profile from an agonist to an antagonist at the M3 and M1 receptors**

We have shown that the presence of acute wash-resistant xanomeline binding results in an increase in basal levels of inositol phosphate production at both the M3 (Chapter 4 Fig. 3) and M1 (Chapter 4 Fig. 4) receptor. However, inositol phosphate levels were no longer elevated following prolonged waiting in control media (Chapter 4 Fig. 8 & Fig. 9). Similar results were obtained in both cases when cells were incubated with xanomeline for 24 hours prior to washing. To determine the time course of the reversal of receptor stimulation by wash-resistant xanomeline binding, cells expressing the M3 receptor were treated with xanomeline for one hour followed by washing and waiting for different durations. The initial receptor activation following pretreatment for one hour and washing subsided rapidly upon further incubation in control media, approaching control levels after 3 hours (Fig. 4 c). As can be seen in Fig. 4 d, a similar response profile was observed when xanomeline was present for increasing durations of time prior to washing (30 minutes to 24 hours), although the time-dependent reversal of persistent receptor activation occurred at a slower rate.

Xanomeline has been shown to antagonize the functional response to carbachol following washing and prolonged incubation in control media at the M3 receptor (Chapter 4 Fig. 8). Further experiments were conducted to determine the time dependence of development of xanomeline antagonistic activity at the M3 receptor. A reduction in the ability of carbachol to elicit a response was observed immediately following xanomeline treatment for one hour. This effect was maintained at a steady state upon further incubation of pretreated and washed cells in control media for various time points up to 23 hours (Fig. 4 c). In contrast, a time-dependent reduction in the ability of carbachol to



stimulate a response was observed when cells expressing the M3 receptor were treated with xanomeline for 30 minutes to 24 hours followed by washing (Fig. 4 d).

Parallel experiments were conducted at the M1 receptor with slightly different results. To determine the time course of the reversal of receptor stimulation by wash-resistant xanomeline, cells expressing the M1 receptor were treated with xanomeline for one hour followed by washing and waiting increasing durations of time. The initial receptor stimulation by wash-resistant xanomeline declined rapidly, reaching control levels after approximately 5 hours of incubation in ligand free media (Fig.4 a). This decrease took slightly longer than it did at the M3 receptor. In contrast, when cells were incubated with xanomeline for increasing amounts of time (30 minute to 24 hours) prior to washing, the initial receptor activation was slightly lower than when xanomeline was continually present in the functional assay. In this case, receptor activation by wash-resistant xanomeline decreased slowly over time, approaching basal levels after 16 hour incubation in control media (Fig. 4 b). This is in stark contrast to the rapid decrease in wash-resistant activation that was observed at the M3 receptor.

Similar to the M3 receptor, xanomeline has also been shown to antagonize the functional response to carbachol following washing and prolonged incubation in control media at the M1 receptor (Chapter 4 Fig. 9). Thus, further experiments were conducted to determine the time dependence of carbachol antagonism by xanomeline. The response to carbachol following xanomeline pretreatment for one hour decreased over time, reaching a maximum of approximately 50 percent inhibition after 9 hours of incubation in control media (Fig. 4 a). However, the decrease in the ability of carbachol to stimulate a functional response following xanomeline pretreatment for 30 minutes to 24 hours decreased in a time-dependent manner in parallel with the decrease in receptor stimulation by wash-resistant xanomeline (Fig. 4 b).

### **Effect of decreasing cell-surface receptor expression using a receptor alkylating agent at the M4 receptor**

The level of receptor expression has been shown to influence binding and functional parameters. The level of receptor expression can change the response to an agonist (Kenakin, 1997). To determine if the lack of long-term regulation of xanomeline at the M2 receptor was due to a lower level of cell-surface receptor density, the level of receptor density was decreased at the M4 receptor to the same levels observed at the M2 receptor. To accomplish this, cells were treated with increasing concentrations of propylbenzyl choline mustard (PrBCM) (0.1 – 10 nM) for 15 minutes followed by washing and the binding of 6.5 nM [<sup>3</sup>H]NMS was determined. There was a concentration dependent decrease in cell-surface receptor density as measured by the decrease in [<sup>3</sup>H]NMS binding, with an approximately 50 percent decrease in binding occurring at 2 nM PrBCM treatment (Fig. 5 a). Thus, 2 nM PrBCM was used in further experiments. When cells expressing M4 receptors were exposed to 2 nM PrBCM for 15 minutes followed by washing and incubation in control media for 24 hours there was no longer a decrease in cell-surface receptor density compared to a no PrBCM treatment control (Fig. 5 b; Table 3). In contrast, when cells were treated with PrBCM for 15 minutes followed by washing and xanomeline treatment for one hour followed by washing and waiting 23 hours in ligand free media there was a decrease in cell-surface receptor density with an increase in radioligand affinity (Fig. 5 b). Similarly, when cells were treated with xanomeline for 24 hours subsequent to treatment with PrBCM there was a decrease in cell-surface receptor density. However, under these pretreatment conditions there was a large decrease in radioligand affinity (Fig. 5 b; Table 3).

### **Effect of blocking the $G_{i/o}$ G proteins on the long-term effects of xanomeline binding at the M4 receptor**

To determine if  $G_{i/o}$  G protein stimulation is necessary for the long-term effects of xanomeline at the M4 receptor cells were pretreated with pertussis toxin (100 ng/ml), which blocks  $G_{i/o}$  G proteins, for 24 hours prior to xanomeline treatments. When cells were treated with xanomeline for one hour followed by washing and further incubation in the absence of free ligand for 23 hours there was decrease in cell-surface receptor density and no change in radioligand affinity compared to control. Similarly, when cells were treated with xanomeline for 24 hours prior to washing there was a decrease in cell-surface receptor expression, however, in this case there was a large decrease in radioligand affinity compared to control (Fig. 6). Overall, inactivation of  $G_{i/o}$  G protein by pertussis toxin did not cause marked changes in receptor regulation by xanomeline (Fig. 6; Chapter 3 Fig. 19 d). Results are summarized in Table 3.

### **Discussion**

Previous research has documented that prolonged exposure to an agonist can lead to receptor internalization, down-regulation or uncoupling from G proteins (Maloteaux & Hermans, 1994). Experiments were therefore conducted to determine whether the observed long-term effects of xanomeline are due to receptor internalization or down-regulation at the M3 and M1 receptors. These possibilities were differentiated by comparing the effects of xanomeline treatment on the specific binding of [ $^3$ H]QNB and [ $^3$ H]NMS (Fig. 1 & Fig. 2). QNB is a lipophilic antagonist that is able to bind to both cell-surface and internalized (but intact) receptors (Ehlert et al., 1981; Vickroy & Malphurs, 1994), whereas NMS (a permanently-charged quaternary amine) is only able to bind to receptors located on the cell surface (Feigenbaum & El-Fakahany, 1985). The lack of

significant differences in the effects of brief xanomeline pretreatment and washing on the binding of both ligands suggests absence of receptor internalization at either receptor subtype. Furthermore, similarities of the effects of 24 hour treatment with xanomeline on the binding profiles of [<sup>3</sup>H]QNB and [<sup>3</sup>H]NMS supports the notion that the effects observed in both displacement and saturation binding experiments are mainly due to receptor down-regulation rather than internalization. In addition, incomplete inhibition of binding of either radioligand across the various xanomeline pretreatments followed by washing at both receptor subtypes suggests that a portion of the cell surface receptors are not susceptible to regulation by xanomeline. Alternatively, this might be an indicator of allosteric modification of receptor conformation by xanomeline, since incomplete inhibition of radioligand binding is a hallmark of allosteric receptor modulation (Lee & El-Fakahany, 1991).

Interestingly, treatment with xanomeline for one hour followed by washing and waiting 23 hours indicated an apparent higher potency of xanomeline in decreasing [<sup>3</sup>H]NMS than [<sup>3</sup>H]QNB binding at the M3 receptor. An opposite profile would have been expected if a major proportion of the receptors were being internalized rather than down-regulated. One possible explanation of these data is that xanomeline wash-resistant binding to the M3 receptor may impart differential allosteric modulation of NMS and QNB binding. The magnitude of cooperativity between the orthosteric and an allosteric site on a receptor is dependent on the specific receptor conformations effected by binding of various ligands (Ehlert, 1988). Therefore, the effects of a given allosteric modulator are ligand specific. In contrast to the changes observed after one hour treatment followed by waiting at the M3 receptor, when cells expressing the M1 receptor were exposed to xanomeline for one minute or one hour followed by washing and waiting there was no

difference in potency of xanomeline in decreasing [<sup>3</sup>H]QNB or [<sup>3</sup>H]NMS binding. This suggests that the observed effects have some degree of receptor specificity.

Additional [<sup>3</sup>H]NMS saturation binding experiments using cells expressing the M3 receptor were conducted at 4°C, which has been shown to prevent receptor activation by agonists (El-Fakahany & Richelson, 1980; Maloteaux & Hermans, 1994). Treatment of cells with xanomeline for one hour at 37°C followed by washing and 23-hour incubation at 4°C did not result in changes in binding parameters compared to untreated control cells incubated at 4°C. However, when cells were treated with 10 or 100 μM xanomeline for 24 hours at 4°C prior to washing there was a decrease in radioligand affinity (Fig. 3). Taken together, these results suggest that xanomeline avidly bound at the receptor allosteric site requires receptor activation for the induction of its long-term effects on cell-surface receptor density. Interestingly, while the effects of xanomeline on maximal cell-surface receptor density seem to be dependent on receptor activation at the orthosteric site, xanomeline's effects on radioligand affinity are not. This differential effect on cell-surface receptor density and radioligand affinity is consistent with other protocols that inhibit receptor function. It has to be noted, however, that changing temperature is known to elicit general changes in the conformation of many cellular proteins in addition to altering receptor conformation.

Increases in activation of phosphoinositide hydrolysis were seen in cells briefly treated with xanomeline and then washed at both the M3 and M1 receptors. However, when pretreated and washed cells were incubated for a long period in the absence of free xanomeline, this wash-resistant receptor activation was reversed to control unstimulated levels. The same phenomenon was observed in cells treated with xanomeline for prolonged durations prior to washing. It is possible that receptor uncoupling from the G protein, internalization or down-regulation could account for these

changes (Cioffi & El-Fakahany, 1989; van de Westerlo et al., 1995). We further explored the time dependence of the development of this biphasic pharmacological profile (Fig. 4). Elevated levels of inositol phosphates that occur during one hour xanomeline treatment followed by washing rapidly decline within the first three hours of incubation in control media in cells expressing the M3 receptor. The profile at the M1 receptor was similar, except the return to basal levels occurred after 5 hour incubation. A similar decline in wash-resistant activation was observed when M3 expressing cells were treated with xanomeline for 30 minutes to 24 hours followed by washing, just prior to performing the receptor functional assay. In contrast, the decline in wash-resistant activation following prolonged treatment with xanomeline occurred more slowly for cells expressing the M1 receptor. The discrepancy between the M3 and M1 receptors may be because xanomeline is a full agonist at the M1 receptor and thus needs fewer functional receptors to still be able to stimulate a response.

Reversal of xanomeline wash-resistant activation of the receptor is accompanied by attenuation of receptor activation by carbachol (Fig. 4). Noteworthy, while there was complete reversal of wash-resistant receptor activation by xanomeline following one hour pretreatment and washing at both receptor subtypes, its antagonistic effects on carbachol reached a maximum of only approximately 25 percent at the M3 receptor. However, the antagonist effect at the M1 receptor was greater reaching approximately a 50 percent decrease in the ability of carbachol to stimulate a response. In contrast, the ability of carbachol to elicit a response following exposure of cells expressing the M3 receptor to xanomeline for 30 minutes to 24 hours followed by washing decreases as the time of xanomeline exposure increases, with a maximal decrease in inositol phosphate production of approximately 67 percent. The response was similar at the M1 receptor; however, the maximal decrease in stimulation was approximately 85 percent. The

higher magnitude of the antagonistic effects of the latter treatment protocol on the response to carbachol might be due to its demonstrated combined effect on receptor number and affinity.

In our experiments, we detected more marked effects of xanomeline on M4 versus M2 receptors. However, the higher level of expression of M4 receptors complicates interpretation of these data. Receptor overexpression has been shown to result in partial agonists appearing to be full agonists in the expression system (Christopoulos & El-Fakahany, 1999; Kenakin, 1997; Wood et al., 1999). Thus, experiments were undertaken to determine if decreasing receptor density using an alkylating agent (PrBCM) at the M4 receptor changes the ability of xanomeline to induce long-term changes. Although receptor availability was decreased by approximately 50 percent prior to xanomeline pretreatments, receptor density in untreated cells had recovered to that of non-alkylated cells 24 hour later. Interestingly, xanomeline pretreatment following treatment with PrBCM still resulted in a decrease in cell-surface receptor density following either long-term xanomeline pretreatment, similar to the decrease observed with xanomeline treatment alone (Chapter 3 Fig. 19). This suggests that an initial decrease in receptor density is not significant in influencing the long-term effects of xanomeline. However, since cell-surface receptor density returned to untreated levels it is possible that the decrease in receptor density did not occur for long enough to detect changes in binding properties due to decreased receptor density.

We have previously demonstrated that long-term receptor regulation at the M1 and M3 receptor subtypes appears to be dependent on receptor function. Blocking function by a mutation in the  $G_{q/11}$  G protein binding domain at the M1 receptor (Chapter 5 Fig. 5) or blocking function using siRNA for  $G_{q/11}$  G proteins at the M3 receptor (Chapter 5 Fig. 7) results in the long-term effects of xanomeline on cell-surface receptor

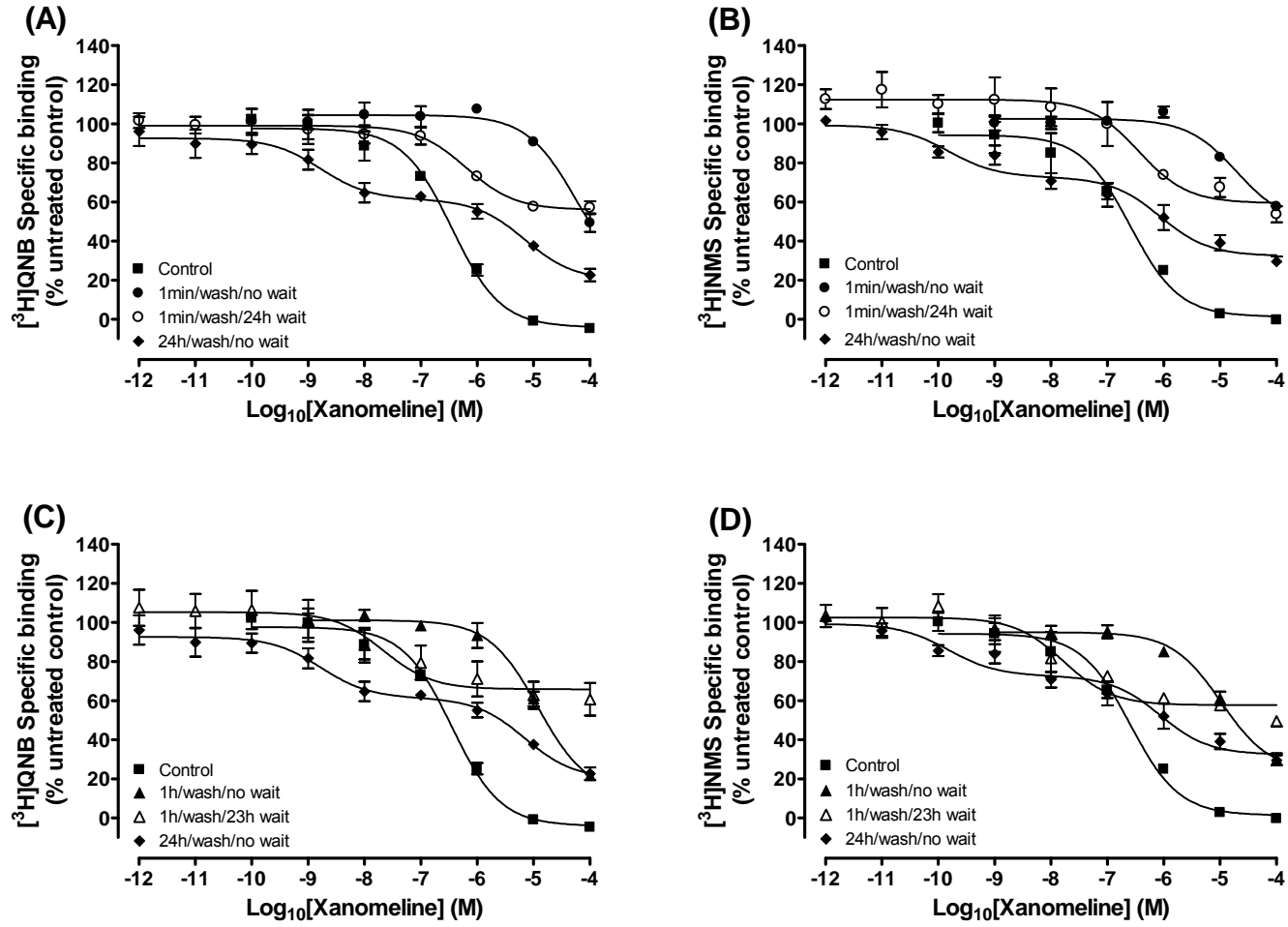
density no longer being apparent. Interestingly, when cells expressing the M4 receptor were treated with pertussis toxin, which blocks the  $G_{i/o}$  G protein (Eglen et al., 1987), there was no change in the binding profile following xanomeline pretreatments compared to xanomeline treatment in the absence of G protein blockade (Fig. 6). This suggests that the long-term regulation of xanomeline at the M4 receptor is mediated by a means other than  $G_{i/o}$  G proteins.

From the combined results of the two previously discussed experiments we propose that the lack of long-term effects of xanomeline at the M2 receptor is due to marked attenuation of receptor response to xanomeline. It is also important to consider that since xanomeline is able to bind, but stimulates a minimal if any functional response, it may have antagonistic properties. Furthermore, this suggests that receptor regulation in response to xanomeline occurs through different mechanisms depending on the receptor subtype.

In summary, prolonged exposure to xanomeline has been shown to lead to changes in receptor regulation. These changes appear to be the result of receptor down-regulation rather than internalization. Long-term receptor regulation is also dependent on a functional response to xanomeline. At both the M1 and M3 receptors this modulation occurs in a time-dependent manner. Furthermore, we have provided evidence to suggest that the long-term modulation of the receptor by xanomeline occurs through different mechanisms at the M1 and M3 receptors compared to the M2 and M4 receptors.

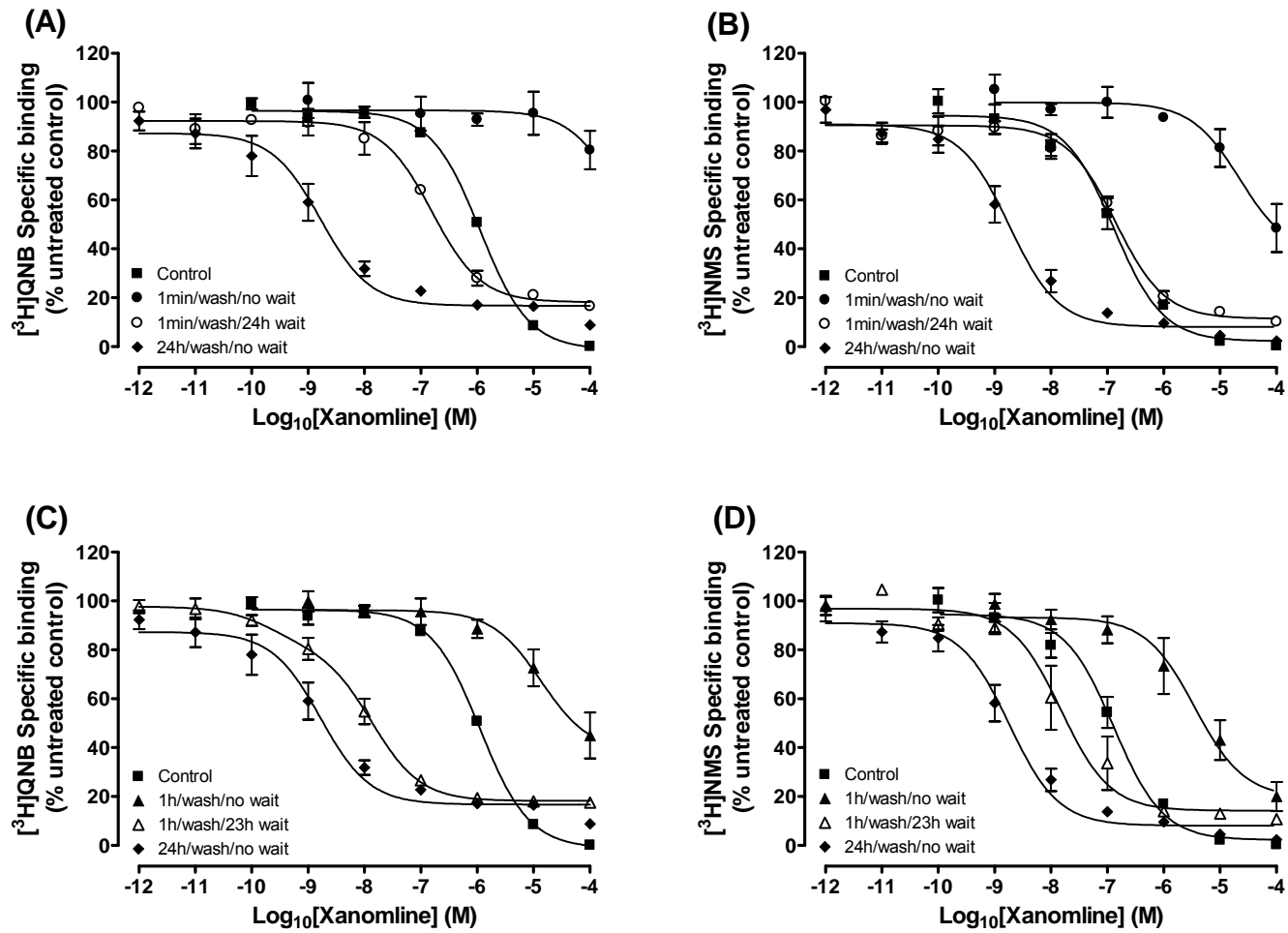


Figure 1



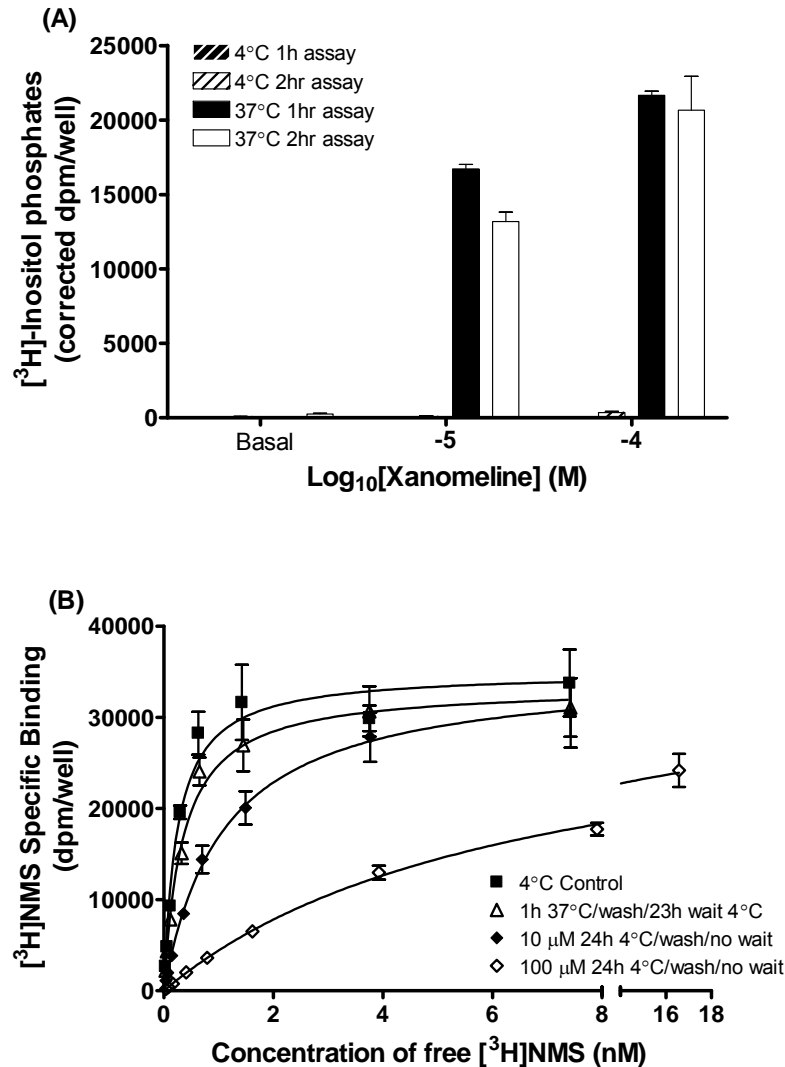
**Figure 1. Comparison of the effects of xanomeline treatment on the specific binding of [<sup>3</sup>H]QNB and [<sup>3</sup>H]NMS in CHO cells expressing hM3 receptors.** A, B) Cells expressing the M3 receptor were pretreated with increasing concentrations of xanomeline for 1 minute followed by washing and immediate use (closed circles) or subsequent to 24 hour incubation in ligand free media (open circles). Additional cells were treated with increasing concentrations of xanomeline for 24 hours prior to washing (closed diamonds). C, D) Cells were pretreated with increasing concentrations of xanomeline for 1 hour, washed and used immediately (closed triangles) or after incubation in control medium for 23 hours (open triangles). An additional group of cells was pretreated with xanomeline for 24 hours prior to washing (closed diamonds). A, C) The binding of 1.4 nM [<sup>3</sup>H]QNB was determined in untreated cells in the concurrent presence of xanomeline (closed squares) or following various xanomeline pretreatments for 1 hour at 37°C. B, D) Cells were treated the same as (A, C) except 2.9 nM [<sup>3</sup>H]NMS was used in place of QNB. Non-specific binding was determined in the presence of 10 μM atropine. Individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 39500 and 56800 dpm/well for QNB and NMS, respectively. Values represent the means ± S.E. of 3-4 experiments conducted in triplicate.

Figure 2



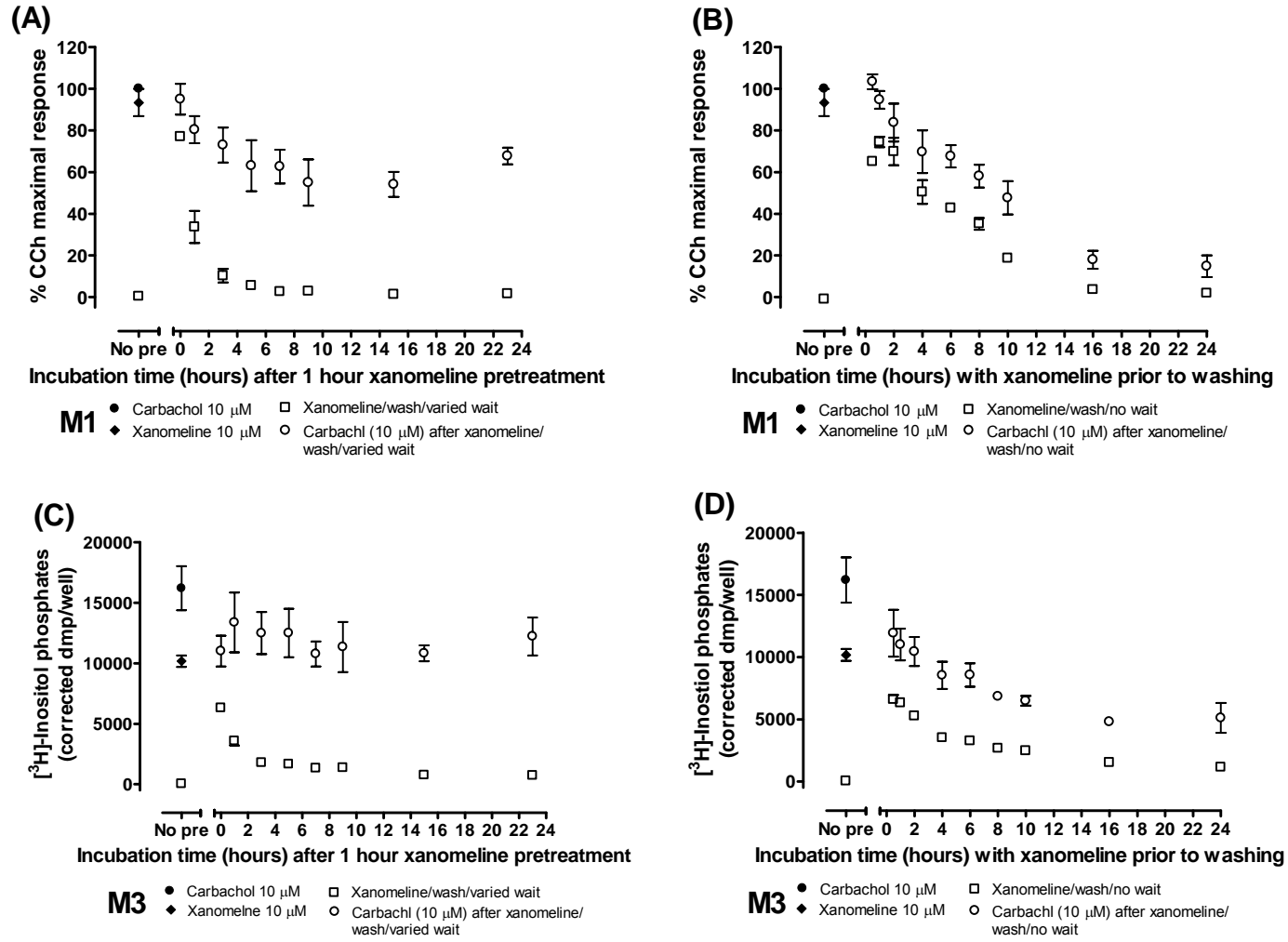
**Figure 2. Comparison of the effects of xanomeline treatment on the specific binding of [<sup>3</sup>H]QNB and [<sup>3</sup>H]NMS in CHO cells expressing hM1 receptors.** A, B) Cells expressing the M1 receptor were pretreated with increasing concentrations of xanomeline for 1 minute followed by washing and immediate use (closed circles) or subsequent to 24 hour incubation in ligand free media (open circles). Additional cells were treated with increasing concentrations of xanomeline for 24 hours prior to washing (closed diamonds). C, D) Cells were pretreated with increasing concentrations of xanomeline for 1 hour, washed and used immediately (closed triangles) or after incubation in control medium for 23 hours (open triangles). An additional group of cells was pretreated with xanomeline for 24 hours prior to washing (closed diamonds). A, C) The binding of 1.4 nM [<sup>3</sup>H]QNB was determined in untreated cells in the concurrent presence of xanomeline (closed squares) or following various xanomeline pretreatments for 1 hour at 37°C. B, D) Cells were treated the same as (A, C) except 2.9 nM [<sup>3</sup>H]NMS was used in place of QNB. Non-specific binding was determined in the presence of 10 μM atropine. Individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 57500 and 80900 dpm/well for QNB and NMS, respectively. Values represent the means ± S.E. of 2-4 experiments conducted in triplicate.

Figure 3



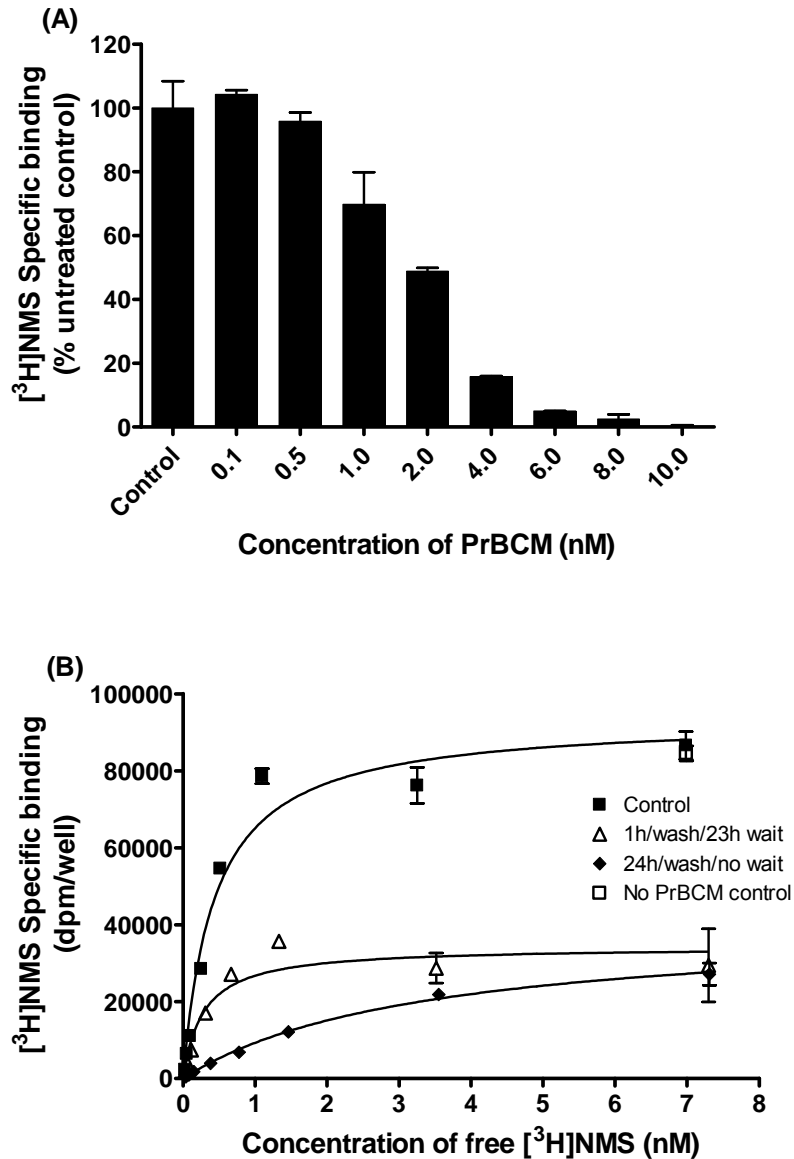
**Figure 3. Effects of blocking receptor activation on the ability of xanomeline to initiate long-term effects on  $[^3\text{H}]\text{NMS}$  saturation binding parameters in CHO cells expressing hM3 receptors.** A) Cells were treated with 10 or 100  $\mu\text{M}$  xanomeline for 1 (black bar) or 2 (white bar) hours in the presence of 10 mM lithium chloride at 4°C (dashed) or 37°C (solid) and the production of inositol phosphates was determined. Values represent 1 experiment conducted in triplicate. B) Cells were pretreated with 10  $\mu\text{M}$  xanomeline for 1 hour at 37°C followed by washing and incubation in the absence of free xanomeline for 23 hours at 4°C (open triangles). Additional groups of cells were treated with 10  $\mu\text{M}$  (closed diamonds) or 100  $\mu\text{M}$  (open diamonds) xanomeline for 24 hours at 4°C prior to washing. Subsequently, untreated cells incubated at 4°C (closed squares) or pretreated cells were incubated with increasing concentrations of  $[^3\text{H}]\text{NMS}$  (0.02 – 16.5 nM) for 1 hour at 37°C. Non-specific binding was determined in the presence of 10  $\mu\text{M}$  atropine. Values represent the means  $\pm$  S.E. of 3-4 experiments conducted in duplicate.

**Figure 4**



**Figure 4. Time course of switching of the pharmacological profile of xanomeline from an agonist to an antagonist in CHO cells expressing hM1 or hM3 receptors.** A) M1 or C) M3 cells were pretreated with 3 or 10  $\mu$ M xanomeline, respectively, for 1 hour followed by washing and waiting for increasing lengths of time (0-23 hours). B) M1 or D) M3 cells were pretreated with 3 or 10  $\mu$ M xanomeline, respectively, for increasing amounts of time (30 minutes -24 hours) prior to washing. Subsequently, the accumulation of inositol phosphates was determined over a 1-hour incubation period at 37°C in the presence of lithium chloride following no further stimulation (open squares) or stimulation with 10  $\mu$ M carbachol (open circles). For comparison, untreated cells were stimulated with 10  $\mu$ M carbachol (closed circles) or 10  $\mu$ M xanomeline (closed diamonds). No pre: untreated cells that were not exposed to xanomeline. Individual experiments at the M1 receptor were normalized to the production of inositol phosphates following 10  $\mu$ M carbachol which had a mean of 31600 and 26700 dpm/well for (A) and (C), respectively. Values represent the means of 2-3 experiments conducted in triplicate.

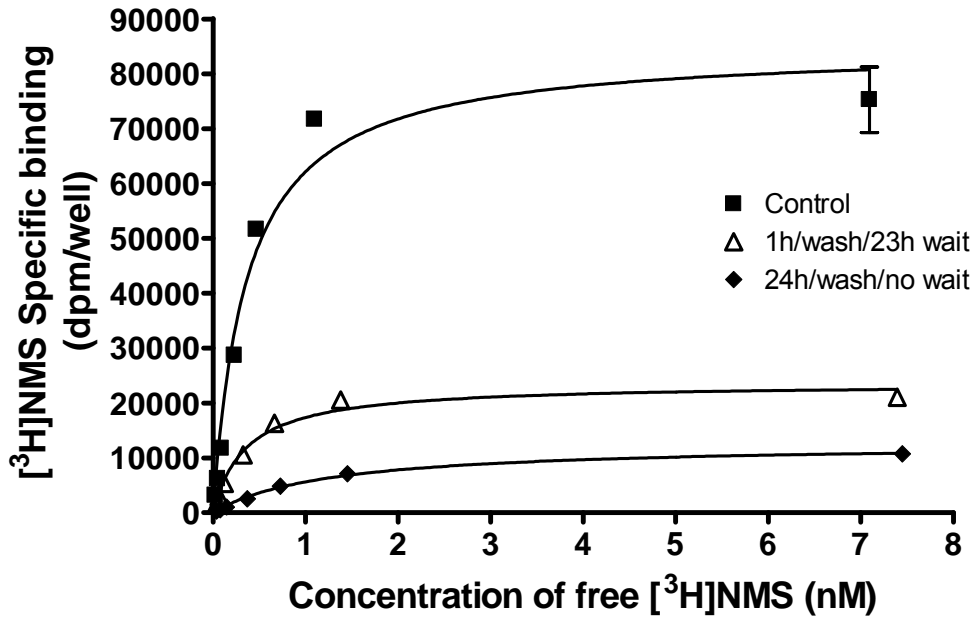
Figure 5



**Figure 5. Effects of decreasing receptor expression using PrBCM on saturation binding parameters in CHO cells expressing hM4 receptors.** A) Cells were pretreated with increasing concentration of cyclized PrBCM (0.1 – 10 nM) for 15 minutes followed by washing. Subsequently, cells were exposed to 6.5 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. B) Cells were pretreated with 2 nM PrBCM prior to xanomeline pretreatments. Cells were pretreated with xanomeline (10 μM) for 1 hour followed by washing and waiting 23 hours (open triangles) or treated with xanomeline for 24 hour prior to washing (closed diamonds). Cells pretreated only with PrBCM (closed squares) or xanomeline pretreated cells were exposed to increasing concentrations of [<sup>3</sup>H]NMS (0.03 – 7 nM) for 1 hour at 37°C. For comparison naïve cells were exposed to the highest concentration of NMS (open square). Non-specific binding was determined in the presence of 10 μM atropine. Values represent 1 experiment conducted in triplicate.



**Figure 6**



**Figure 6. Effects of blocking the  $G_{i/o}$  G proteins on the ability of xanomeline to induce long-term changes in CHO cells expressing the hM4 receptor.** Cells were pretreated with 100 ng/ml pertussis toxin for 24 hours. Subsequently, cells were pretreated with 10  $\mu$ M xanomeline for 1 hour followed by washing and 23 hour incubation in control media containing pertussis toxin (open triangles). An additional group of cells was treated concurrently with xanomeline and pertussis toxin for 24 hours prior to washing (closed diamonds). Untreated (closed squares) or xanomeline pretreated cells were then exposed to increasing concentrations of [ $^3$ H]NMS (0.02 – 7.5 nM) for 1 hour at 37°C. Non-specific binding was determined in the presence of 10  $\mu$ M atropine. Values represent 1 experiment conducted in triplicate.

**Table 1. Inhibition of radioligand binding using a receptor-saturating concentration of [<sup>3</sup>H]QNB or [<sup>3</sup>H]NMS following xanomeline pretreatment and various waiting periods in CHO cells expressing hM1 or hM3 receptor subtypes.** Cells were treated with increasing concentrations of xanomeline (1 pM – 100 μM) for 1 minute or 1 hour followed by washing and immediate use in the binding assay or subsequent to incubation for 24 or 23 hours, respectively, in the absence of free xanomeline. An additional group of cells was treated with xanomeline for 24 hours prior to washing. Subsequently, untreated cells were incubated concurrently with increasing concentration of xanomeline and 1.4 nM [<sup>3</sup>H]QNB or 2.9 nM [<sup>3</sup>H]NMS or pretreated cells were incubated with 1.4 nM [<sup>3</sup>H]QNB or 2.9 nM [<sup>3</sup>H]NMS for 1 hour at 37°C. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

Pretreatment group	<u>[<sup>3</sup>H]QNB</u>				<u>[<sup>3</sup>H]NMS</u>			
	pIC <sub>50</sub> <sup>a</sup>	pIC <sub>50-H</sub> <sup>b</sup>	pIC <sub>50-L</sub> <sup>c</sup>	I <sub>max</sub> <sup>d</sup>	pIC <sub>50</sub>	pIC <sub>50-H</sub>	pIC <sub>50-L</sub>	I <sub>max</sub>
<b>M1 receptor</b>								
Control (untreated) <sup>e</sup>	5.9 ± 0.03			102 ± 1.5	6.9 ± 0.08			98 ± 1.0
1min/washout/no wait	3.7 ± 0.57 <sup>~</sup>			ND	4.6 ± 0.04 <sup>&amp;</sup>			63 ± 10.3 <sup>‡</sup>
1min/washout/24h wait	6.8 ± 0.04 <sup>‡</sup>			82 ± 0.2	6.9 ± 0.03 <sup>@</sup>			89 ± 0.9
1 h/washout/no wait	4.8 ± 0.08 <sup>#</sup>			62 ± 9.6 <sup>‡</sup>	5.5 ± 0.20 <sup>\$</sup>			80 ± 3.6 <sup>‡</sup>
1 h/washout/23 h wait	8.3 ± 0.19 <sup>†</sup>			74 ± 2.0 <sup>‡</sup>	7.8 ± 0.38 <sup>*</sup>			84 ± 07 <sup>‡</sup>
24 h/washout/no wait	9.2 ± 0.30 <sup>+†</sup>			72 ± 2.2 <sup>‡</sup>	8.8 ± 0.26 <sup>@*</sup>			88 ± 0.7 <sup>‡</sup>
<b>M3 receptor</b>								
Control (untreated)	6.4 ± 0.11			104 ± 0.5	6.6 ± 0.16			97 ± 1.2
1min/washout/no wait	4.2 ± 0.06 <sup>~</sup>			82 ± 7.5 <sup>‡</sup>	4.7 ± 0.16 <sup>&amp;</sup>			53 ± 5.9 <sup>‡</sup>
1min/washout/24h wait	6.2 ± 0.19 <sup>‡</sup>			44 ± 3.2 <sup>‡</sup>	6.5 ± 0.08 <sup>@</sup>			40 ± 3.0 <sup>‡</sup>
1 h/washout/no wait	4.9 ± 0.12 <sup>#</sup>			88 ± 2.6	5.0 ± 0.09 <sup>\$</sup>			77 ± 1.9 <sup>‡</sup>
1 h/washout/23 h wait	6.7 ± 0.18 <sup>†</sup>			34 ± 8.0 <sup>‡</sup>	7.9 ± 0.30 <sup>*</sup>			43 ± 1.1 <sup>‡</sup>
24 h/washout/no wait		8.8 ± 0.13 <sup>+†</sup> (43 ± 3.3%)	5.1 ± 0.01	80 ± 3.2 <sup>‡</sup>		9.2 ± 0.47 <sup>@*</sup> (42 ± 3.3%)	6.0 ± 0.37	70 ± 1.5 <sup>‡</sup>

Table 1 cont.

<sup>a</sup> Negative logarithm of the IC<sub>50</sub> for binding to a single potency site.

<sup>b</sup> Negative logarithm of the IC<sub>50</sub> for the high-potency agonist binding site; percent of binding sites shown in parentheses.

<sup>c</sup> Negative logarithm of the IC<sub>50</sub> for the low-potency agonist binding site.

<sup>d</sup> Maximal percent inhibition of radioligand binding.

<sup>e</sup> Control cells were incubated simultaneously with xanomeline and the radioligand.

~ Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of QNB binding between the 1min/wash/no wait group compared to control.

± ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of QNB binding where indicated between the 1min/wash/24h wait or 24h/wash/no wait pretreated groups compared to xanomeline 1min/wash/no wait treatment.

# Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of QNB binding between the 1h/wash/no wait group compared to control.

† ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of QNB binding where indicated between the 1h/wash/23h wait or 24h/wash/no wait pretreated groups compared to xanomeline 1h/wash/no wait treatment.

& Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of NMS binding between the 1min/wash/no wait group compared to control.

@ ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of NMS binding where indicated between the 1min/wash/24h wait or 24h/wash/no wait groups compared to xanomeline 1min/wash/no wait treatment.

\$ Unpaired t-test detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of NMS binding between the 1h/wash/no wait group compared to control.

\* ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in pIC<sub>50</sub> of inhibition of NMS binding where indicated between the 1h/wash/23h wait or 24h/wash/no wait groups compared to xanomeline 1h/wash/no wait treatment.

‡ Significantly different ( $p < 0.05$ ) from 100 percent as determined by ANOVA followed by Dunnett's post-test comparison.

**Table 2. Effects of blocking receptor activation on xanomeline-induced changes in [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing hM3 receptors.** Cells were treated with 10 μM xanomeline for 1 hour at 37°C followed by washing and used after 23 hour incubation in control media at 4°C. Additional cells were treated with 10 or 100 μM xanomeline for 24 hours at 4°C prior to washing. Untreated control cells were incubated at 4°C for 24 hours. Subsequently, untreated or treated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.03 – 16.6) at 37°C for 1 hour. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

<b>Xanomeline pretreatment group</b>	<b>K<sub>d</sub><sup>a</sup> (nM)</b>	<b>B<sub>max</sub><sup>b</sup></b>
Control (untreated) 4°C	0.24 ± 0.004	35000 ± 2800
1 h 37°C /washout/23 h 4°C wait	0.34 ± 0.03	33400 ± 3600
24 h 4°C (10 μM)/washout/no wait	1.08 ± 0.06*	35300 ± 4100
24 h 4°C (100 μM)/washout/no wait	6.25 ± 0.44*	32500 ± 2000

<sup>a</sup> Equilibrium dissociation constant for [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell-surface receptor density (dpm/well).

\* ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in K<sub>d</sub> or B<sub>max</sub> where indicated between the pretreated groups compared with control.

**Table 3. Effects of changing the level of receptor expression or inactivating the G proteins on xanomeline-induced changes in [<sup>3</sup>H]NMS saturation binding parameters in CHO cells expressing the hM4 receptor subtype.** Cells were pretreated with PrBCM (2 nM) for 15 minutes followed by washing and then subjected to xanomeline pretreatments. Additional cells were treated with pertussis toxin (100 ng/ml) for 24 hours prior to xanomeline pretreatments as noted in methods. Xanomeline treatment: cells were treated with 10 μM xanomeline for 1 hour followed by washing and incubation for 23 hours in the absence of free ligand. An additional group of cells was treated with xanomeline for 24 hours prior to washing. Control untreated or xanomeline pretreated cells were then incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.01 – 7.5 nM) for 1 hour at 37°C. Parameters are from a single experiment.

	<u>PrBCM</u>		<u>Pertussis toxin</u>	
	<b>K<sub>d</sub></b> <sup>a</sup>	<b>B<sub>max</sub></b> <sup>b</sup>	<b>K<sub>d</sub></b>	<b>B<sub>max</sub></b>
Control (untreated)	0.44	93600	0.37	84900
1 h/washout/23h wait	0.28	34300	0.36	23600
24 h/washout/no wait	3.32	40200	1.26	12700

<sup>a</sup> Equilibrium dissociation constant for [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell surface receptor density (dpm/well).

Statistical analyses were not performed because the data presented are from single experiments conducted in triplicate.

## **CHAPTER 7: Conclusions**

## **Conclusions**

Muscarinic receptors are potential targets for the development of drugs to treat diseases of the central nervous system such as Alzheimer's disease and schizophrenia. Currently, cholinergic drug treatment has shown promise in reducing symptoms such as cognitive deficits, hallucinations and delusions, which are thought to be mediated by the M1 and perhaps M4 muscarinic receptors. However, there are negative side-effects observed with cholinergic drugs that are thought to be mediated by non-selective activation of M2, M3 and M5 muscarinic receptor subtypes. A major goal of current research in muscarinic receptor biology and pharmacology is to develop agonists and antagonists that act in a subtype-selective manner, potentially through binding to allosteric sites on the receptor that are less conserved among subtypes. In order to accomplish this, we need to learn more about the allosteric site(s) on the muscarinic receptors and how potential drugs interact with these regions to modify the action of the receptor.

My thesis addressed a small part of this goal by focusing on the consequences of the long-term interaction of the novel muscarinic receptor agonist xanomeline with M1-M4 subtypes of muscarinic receptors, with a more detailed focus on the mechanisms of action of xanomeline at the M1 and M3 subtypes of muscarinic receptors. Xanomeline has been shown to act as an allosteric modulator of at least the M1 muscarinic receptor. More specifically, the questions that were asked were 1) does xanomeline bind to all receptor subtypes in a reversible and wash-resistant manner?, 2) what are the short- and long-term consequences of xanomeline binding on function at the M1 and M3 receptor subtypes?, 3) what is the role of the orthosteric binding site in the long-term effects of xanomeline at the M1 and M3 receptor subtypes? and 4) what is the

mechanism of receptor regulation as a result of the long-term consequences of xanomeline binding?

My research demonstrated that xanomeline is able to bind in a reversible manner at the orthosteric site and in a wash-resistant manner at an allosteric site at all of the subtypes tested. Interestingly, the potency of xanomeline binding across the four subtypes was similar when xanomeline was present acutely. However, differences in the binding profiles were apparent between subtypes when xanomeline was present for prolonged periods of time. These differences were most striking at the M2 receptor. However, there were also subtle differences between the other subtypes with the M1 and M3 receptors having the most similar binding profiles. My research further supports that xanomeline interacts with muscarinic receptors in a manner that is unique from other conventional muscarinic agonists and that there is a time-dependent component associated with that interaction. These results are important because it gives some insight into the consequences of prolonged binding of an agonist at the allosteric site.

The unique binding properties of xanomeline led us to determine the functional consequences of xanomeline exposure at the M1 and M3 receptor subtypes. Xanomeline was shown to be a full agonist at the M1 receptor and a partial agonist at the M3 receptor. Wash-resistant binding of xanomeline is able to stimulate a functional response at both receptor subtypes. Both acute and long-term treatments with xanomeline modulate the ability of additional agonists to stimulate a response. The major difference that is observed as a result of xanomeline treatment between the two subtypes is that the presence of xanomeline antagonizes the response to the subsequent addition of agonists to a greater extent at the M3 receptor than at the M1 receptor. These results demonstrate that the minor differences observed in binding studies are mirrored by minor differences in effects on receptor function. Furthermore,



these results are important because they demonstrate the ability of xanomeline to modulate the function of the receptor through binding at an allosteric site.

However, the long-term effects of xanomeline exposure are quite similar at the M1 and M3 muscarinic receptors in terms of cell-surface receptor density and radioligand binding affinity. In both cases activation of the orthosteric site is required to obtain the effects of long-term receptor regulation by xanomeline. This is of interest because xanomeline has been found to be a full agonist at the M1 receptor, but a partial agonist at the M3 receptor. This suggests that although there may be some degree of functional subtype selectivity initially, there is relatively little selectivity in the long-term effects of xanomeline exposure between these two subtypes. This is a potential problem for using xanomeline to selectively target M1 receptors. This is also important because it emphasizes that when drugs are being screened it is not only important to determine the properties of the initial exposure, but the consequences of prolonged exposure also need to be considered.

Furthermore, my research demonstrates that the mechanism of receptor regulation following prolonged exposure to xanomeline is most likely via down-regulation of the receptor. Again, xanomeline exposure appears to result in receptor modulation that is similar between the M1 and M3 receptor subtypes, although the time dependence of this is slightly different. These results are similar to what is observed following prolonged exposure of muscarinic receptors to conventional muscarinic receptor agonists, further supporting the role of the orthosteric binding site in the long-term effects of xanomeline. Interestingly, it appears that the mechanism of receptor regulation is different between the M2 and M4 receptor subtypes compared to the M1 and M3 receptor subtypes. Blocking G protein function at the M4 receptor had no effect on the long-term consequences of xanomeline exposure; however, blocking signaling through

the G protein at the M1 and M3 receptors abolished the long-term effects of xanomeline. These results demonstrate that the nature of drug-receptor interactions is not only dependent upon the type of binding, but also the signaling mechanisms that are activated subsequent to binding.

All of the experiments that I presented were conducted in an *in vitro* receptor expression system; individual receptors expressed in CHO cells. This approach has its strengths and drawbacks. The advantage of using this type of experimental set up is that only a single subtype of receptor is expressed so the actions of xanomeline at a particular subtype can be determined. This would not be possible using tissue preparations since tissues are known to express multiple subtypes of muscarinic receptors. However, there are potential problems associated with the use of receptor expression systems such as there may be differences in the coupling of G proteins and other associated proteins than would be observed endogenously, as well as receptor overexpression could result in changes in binding and functional properties of xanomeline. Finally, it is possible that the concentrations of xanomeline tested in these experiments may be higher than the concentrations that would be found at the target site(s) *in vivo*. Thus, further experimentation *in vivo* is necessary to confirm the results that we obtained in our system and further elucidate the effects of xanomeline treatment.

In addition to conducting further experiments *in vivo*, there are other questions that still remain that would be more easily addressed *in vitro*. Some of these questions include, Where is the allosteric site that xanomeline binds to located? Is the allosteric site located in similar regions across receptor subtypes? What residues are important in the development of wash-resistant binding? Is the allosteric site unique to xanomeline or do other allosteric modulators utilize the same site? What is the signaling pathway/mechanism that is activated as a result of xanomeline stimulation? Are there

any interactions/what are the interactions between receptor subtypes when exposed to xanomeline? These questions can be addressed using a variety of different methods. For example, studies utilizing receptor mutagenesis can be employed to determine the location of the allosteric binding pocket and the residues important in the formation of wash-resistant binding. The creation of receptor chimeras could address questions about the signaling pathways that are involved. Furthermore, expression of multiple subtypes of receptors in the same cell would be useful in determining the potential interaction between different subtypes of muscarinic receptors following drug exposure. It would also be important to extend these studies to look at other allosteric modulators of muscarinic receptors to determine if there is a common mechanism of action between allosteric modulators or if each modulator acts in its own specific manner. Together the answers to these questions would give us more insight and understanding of the mechanisms of action of allosteric modulators of muscarinic receptors. This information would aid in the development of agonists and antagonists that act in a subtype selective manner.

In summary, my data demonstrate that the novel muscarinic receptor agonist xanomeline interacts with muscarinic receptors in a unique manner. Although the acute effects of xanomeline exposure are similar across subtypes the long-term effects are somewhat distinctive between the receptor subtypes. The binding and functional characteristics of xanomeline confirm that it is unlikely to be a beneficial therapeutic due to its interaction with multiple muscarinic receptor subtypes. Thus, my research demonstrates that it is important to determine the long-term interactions of drugs on receptor properties in addition to the acute effects as part of the drug discovery and development process. This is an exciting time for the field of muscarinic receptor biology. A plethora of new research is directed at understanding and developing

allosteric modulators of muscarinic receptors in the hope that they will display subtype selectivity and thus can be used in the treatment of various diseases of the central nervous system.

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**APPENDIX A: Differences observed between suspension and monolayer protocols**



## **Introduction**

The results of an experiment can be dependent on the protocol used to address the question. For example, previous research by Thompson & Fisher (1990) found that treating cells in monolayer with agonists resulted in a larger decrease in receptor expression levels compared to the treatment of cells in suspension. In another case, when forskolin-stimulated cAMP accumulation was studied in cells in suspension it only initiated 75 percent of the response that was observed in cells assayed in monolayer (Mistry et al., 2005). However, the opposite was true in another assay conducted by Mistry et al. (2005). In that case they found that pilocarpine elicited a larger response when cells were stimulated in suspension rather than in monolayer. Together, these results demonstrate that the response to agonists is dependent on both the assay conditions and the agonist being studied.

In our experimental protocols we noted some differences following xanomeline treatment depending on whether the assay was conducted on cells in monolayer or in suspension. To address this question we performed experiments utilizing various binding paradigms and compared the results between the protocols. Our data demonstrate that there are some differences between monolayer and suspension binding protocols following xanomeline treatments, so when possible experiments were conducted on cells in monolayer at all receptor subtypes.

## **Results**

### **Comparison between saturation binding with xanomeline pretreatments conducted in monolayer versus in suspension in hM3 CHO cells**

The binding profile of xanomeline was determined using two different protocols. In either case, pretreatment with 10  $\mu$ M xanomeline for one hour was conducted on cells

in monolayer. Subsequently, cells were exposed to increasing concentrations of [<sup>3</sup>H]NMS either in suspension or monolayer. Pretreatment with xanomeline for one hour followed by washing and immediate use in the suspension binding assay resulted in no change in cell-surface receptor density but a slight decrease in radioligand affinity compared to control; however, it did not reach significance (Fig. 1 a). In contrast, as presented in chapter 3, when the same pretreatment conditions were used but the assay was conducted in monolayer there was a significant ( $p < 0.05$ ) decrease in radioligand affinity (Fig. 1 b). When cells were treated with xanomeline for one hour followed by washing and waiting 23 hours there was no change in radioligand affinity and only a slight decrease in cell-surface receptor density for cells that were assayed in suspension (Fig. 1 a). In contrast, the same pretreatments followed by assaying in monolayer resulted in a significant ( $p < 0.05$ ) decrease in cell-surface receptor expression (Fig. 1 b). When pretreatment time with xanomeline was increased to 24 hours prior to washing and performing the binding assay in suspension there was a significant ( $p < 0.05$ ) decrease in radioligand affinity, however, there was only a slight decrease in cell-surface receptor density (Fig. 1 a). In contrast, when the assay was conducted in monolayer there was a significant ( $p < 0.05$ ) decrease in both parameters (Fig. 1 b). Results are summarized in Table 1.

### **Comparison between the shift in inhibition of [<sup>3</sup>H]NMS binding curves following acute xanomeline exposure in monolayer versus in suspension at the hM1 CHO cells**

Experiments were conducted to determine the inhibition of [<sup>3</sup>H]NMS binding as a result of exposure of cells expressing the M1 receptor to xanomeline for one hour followed by washing and immediate use in the binding assay. These results were

compared to the ability of xanomeline to inhibit [<sup>3</sup>H]NMS binding when it was present during the binding assay. When cells were assayed in suspension there was a decrease in potency of 1.8 orders of magnitude when xanomeline was present for one hour followed by washing (pEC<sub>50</sub> 5.6) compared to when xanomeline was continually present (control; pEC<sub>50</sub> 7.4) in the binding assay (Fig. 2 a). In contrast, when cells were assayed in monolayer the decrease in potency was only 0.5 orders of magnitude for xanomeline pretreatment (pEC<sub>50</sub> 5.7) compared to control (pEC<sub>50</sub> 6.2) (Fig. 2 b).

## **Discussion**

Alterations in experimental protocols have previously been shown to result in changes in the binding and functional profiles of receptors. These changes appear to be a result of both the protocol employed and the agonist studied. Experiments were therefore conducted to determine if there were differences in the binding properties following exposure to xanomeline when cells were assayed in monolayer compared to suspension. Our results show that pretreatment with xanomeline resulted in a larger decrease in cell-surface receptor expression when cells expressing the M3 receptor were assayed in monolayer compared to suspension (Fig. 1). Additionally, the shift in potency between control and xanomeline treatment for one hour followed by washing and immediate use was larger for cells expressing the M1 receptor assayed in suspension compared to monolayer (Fig. 2).

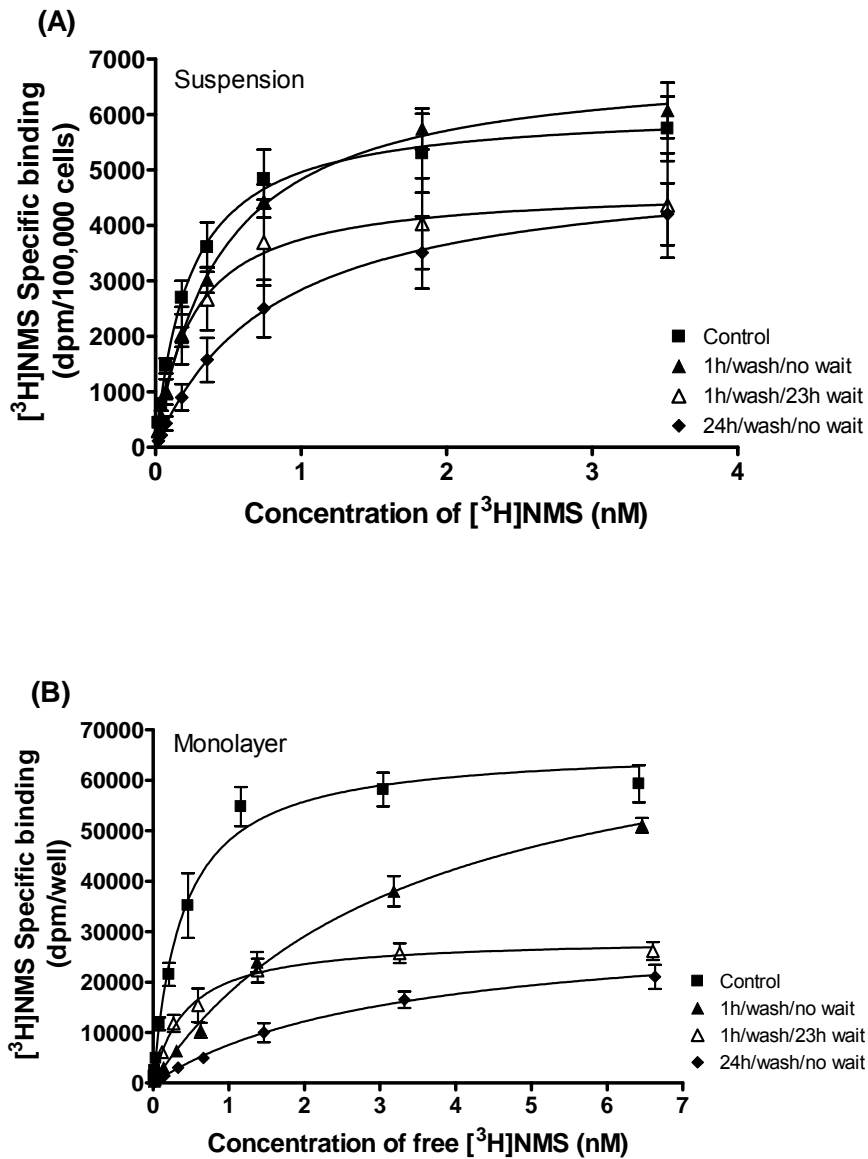
The cause of the difference in binding profiles between the different assays was not studied in detail. However, a potential explanation could be that the differences are a result of differential effects of cytoskeletal elements in cells that are floating (suspension) versus attached (monolayer). Receptor activation has been shown to result in changes in cytoskeletal arrangement (Grimes & Miettinen, 2003). Furthermore,

endocytosis is thought to be dependent on recruitment of beta-arrestin, which is a scaffolding protein (Grimes & Miettinen, 2003). Additionally, it is possible that the procedure employed to detach the cells prior to the binding assay alters the binding profile of xanomeline that is bound in a wash-resistant manner.

Interestingly, the levels of protein for the saturation experiments with cells assayed in suspension were one tenth of the protein levels seen in monolayer. This difference corresponds to the ten fold difference in maximal cell-surface receptor expression levels observed between suspension and monolayer. Thus, the difference in saturation binding profiles between the two methods is most likely not a result of differences in receptor expression.

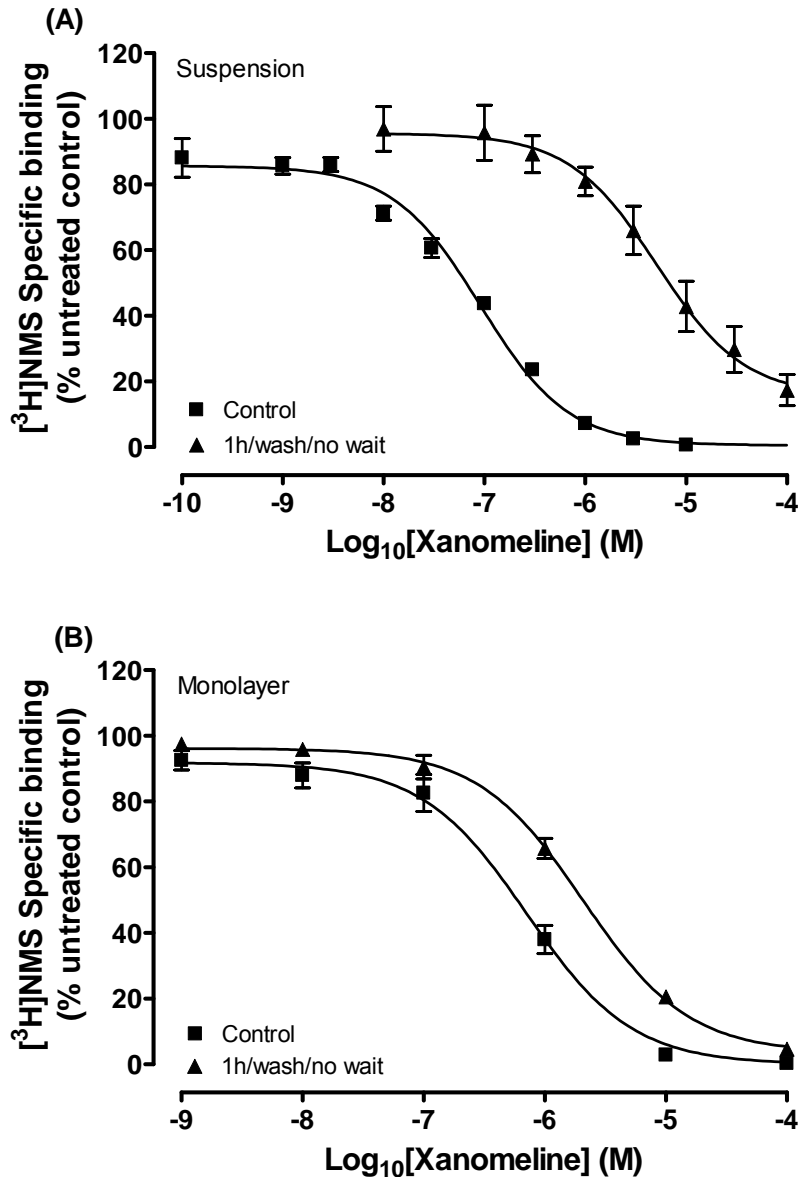
In summary, we addressed the question, is there a difference in the binding properties following exposure to xanomeline at either or both the M1 and M3 receptors when cells are assayed in suspension compared to monolayer? In both cases we observed slight differences in the binding properties suggesting that the difference is dependent on the protocol and not the specific receptor subtype studied. However, this was by no means an extensive study of the differences between the binding properties of cells assayed in monolayer compared to suspension, but rather an observation that was made early on that directed the protocols employed in further studies. Thus, when possible experimental assays were performed on cells in monolayer at all receptor subtypes so direct comparisons between groups could be made.

Figure 1



**Figure 1. Comparison between [<sup>3</sup>H]NMS saturation binding paradigms on assays conducted in suspension versus in monolayer in CHO cells stably expressing hM3 receptors.** Cells were pretreated with 10  $\mu$ M xanomeline for 1 hour followed by washing and used immediately (closed triangles) or subsequent to 23 hour incubation in ligand free media (open triangles). Additional cells were treated with xanomeline for 24 hours prior to washing (closed diamonds). Subsequently, untreated (closed squares) or xanomeline pretreated cells were incubated with increasing concentrations of [<sup>3</sup>H]NMS (0.01 –6.6 nM) for 1 hour at 37°C. The radioligand binding assay was conducted in A) suspension or B) monolayer. Values represent the means  $\pm$  S.E. of 4 experiments conducted in triplicate.

Figure 2



**Figure 2. Comparison between inhibition of  $[^3\text{H}]\text{NMS}$  binding following xanomeline pretreatments for assays conducted in suspension versus in monolayer in CHO cells stably expressing hM1 receptors.** Cells were pretreated with increasing concentrations of xanomeline for 1 hour followed by washing and used immediately (closed triangles). Subsequently, cells were incubated with 0.2 nM  $[^3\text{H}]\text{NMS}$  concurrently with xanomeline (closed squares) or following xanomeline pretreatments for 1 hour at 37°C. The radioligand binding assay was conducted in A) suspension or B) monolayer. For (B) individual experiments were normalized to control binding in the absence of xanomeline treatment which had a mean of 13200 dpm/well. Values represent the means  $\pm$  S.E. of 3-6 experiments conducted in triplicate. Data in (A) provided by Marianne Grant.

**Table 1. Effects on [<sup>3</sup>H]NMS saturation binding parameters when CHO hM3 cells were assayed in monolayer versus suspension.** Cells were treated with 10 μM xanomeline for 1 hour followed by washing and immediate use in the binding assay or subsequent to incubation for 23 hours in the absence of free agonist. An additional group of cells was treated with agonist for 24 hours prior to washing. Control untreated or agonist-pretreated cells were then incubated either in monolayer or in suspension with increasing concentrations of [<sup>3</sup>H]NMS for 1 hour at 37°C. Parameters (± S.E.M.) were derived from computer-assisted non-linear regression analysis.

Xanomeline pretreatment group	<u>Monolayer</u> <sup>†</sup>		<u>Suspension</u>	
	K <sub>d</sub> <sup>a</sup> (nM)	B <sub>max</sub> <sup>b</sup>	K <sub>d</sub> (nM)	B <sub>max</sub>
Control (untreated)	0.42 ± 0.06	67700 ± 3200	0.23 ± 0.01	6100 ± 720
1 h/washout/no wait	3.52 ± 0.52*	78500 ± 1600	0.45 ± 0.01	7000 ± 530
1 h/washout/23 h wait	0.59 ± 0.21	30400 ± 2000*	0.26 ± 0.01	4700 ± 970
24 h/washout/no wait	4.19 ± 0.93*	37000 ± 4600*	1.10 ± 0.32*	5400 ± 610

<sup>a</sup> Equilibrium dissociation constant of [<sup>3</sup>H]NMS binding.

<sup>b</sup> Cell-surface receptor density (dpm/well).

<sup>†</sup> Values are from saturation binding experiments presented in chapter 3.

\*ANOVA followed by Dunnett's post-test comparison detected a significant difference ( $p < 0.05$ ) in K<sub>d</sub> or B<sub>max</sub> between the xanomeline pretreated groups compared with their untreated control.