

# Identification and Characterization of Multiple Subtypes of Muscarinic Acetylcholine Receptors and Their Physiological Functions in Canine Hearts

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## ABSTRACT

M<sub>2</sub> receptors have long been believed to be the only functional subtype of muscarinic acetylcholine receptor (mAChR) in the heart, although recent studies have provided evidence for the presence of other subtypes. We performed a detailed study to clarify this issue. In the presence of tetramethylammonium (1 μM to 10 mM), a novel K<sup>+</sup> current with both delayed rectifying and inward rectifying properties (I<sub>KTMA</sub>) was activated in single canine atrial myocytes. 4-Aminopyridine (0.05–2 mM) also induced a K<sup>+</sup> current (I<sub>K4AP</sub>) with characteristics similar to but distinct from those of I<sub>KTMA</sub>. Both I<sub>KTMA</sub> and I<sub>K4AP</sub> were abolished by 1 μM atropine. I<sub>K4AP</sub>, but not I<sub>KTMA</sub>, was minimized by treatment with pertussis toxin. I<sub>KTMA</sub> was markedly decreased by 4-diphenylacetoxy-N-methylpiperidine methiodide (a selective antagonist for M<sub>3</sub> subtype) but was not altered by pirenzepine (for M<sub>1</sub>), methoctramine (for M<sub>2</sub>), and tropicamide (for

M<sub>4</sub>). Tropicamide substantially reduced I<sub>K4AP</sub>, but the antagonists for other mAChR subtypes had no effects on I<sub>K4AP</sub>. By comparison, I<sub>KACH</sub> (ACh-induced K<sup>+</sup> current) was significantly depressed by methoctramine but was unaltered by other antagonists. Results from displacement binding of [methyl-<sup>3</sup>H]N-scopolamine methyl chloride with pirenzepine, methoctramine, 4-diphenylacetoxy-N-methylpiperidine methiodide, or tropicamide revealed the coexistence of multiple mAChR subtypes in canine atrium. Cloning of cDNA fragments and detection of mRNAs coding for M<sub>2</sub>, M<sub>3</sub>, and M<sub>4</sub> provided further supporting evidence. Our results suggest that 1) multiple subtypes of mAChRs (M<sub>2</sub>/M<sub>3</sub>/M<sub>4</sub>) coexist in the dog heart and 2) different subtypes of mAChRs are coupled to different K<sup>+</sup> channels. Our findings represent the first functional evidence for the physiological role of cardiac M<sub>3</sub> and M<sub>4</sub> receptors.

Accumulating evidence indicates that the functional responses of cells to muscarinic acetylcholine receptors (mAChRs) are mediated by multiple receptor subtypes (Hulme et al., 1990; van Zwieten and Doods, 1995). The heterogeneity of mAChR was first suggested on the basis of pharmacological profiles (differential binding affinity and functional studies) of receptor agonists and antagonists (Goyal, 1989; Hulme et al., 1990). To date, at least four different subtypes have been pharmacologically and functionally defined in primary tissues, designated M<sub>1</sub>, M<sub>2</sub>, M<sub>3</sub>, and M<sub>4</sub> (Eglen and Whiting, 1990; Mutschler et al., 1995; van Zwieten and Doods, 1995). Five isoforms of mAChR have been identified based on molecular cloning studies, desig-

nated m<sub>1</sub> through m<sub>5</sub>, each of which is encoded by an intronless gene with unique amino acid sequence and distinct tissue distribution (Bonner et al., 1987; Peralta et al., 1987; Goyal, 1989). Four of these cloned subtypes (m<sub>1</sub>, m<sub>2</sub>, m<sub>3</sub>, and m<sub>4</sub>) correspond to the functional receptors M<sub>1</sub>, M<sub>2</sub>, M<sub>3</sub>, and M<sub>4</sub>, whereas the physiological relevance of m<sub>5</sub> remains to be established.

The M<sub>2</sub> receptor is commonly believed to be the only functional mAChR subtype in the heart (Watson et al., 1983; Bonner et al., 1987; Mizushima et al., 1987; Peralta et al., 1987) and contributes to the regulation of the heart rate and contractility and to shaping of the action potentials (Jeck et al., 1988; van Zwieten and Doods, 1995). Binding of acetylcholine (ACh) to mAChRs in cardiac cells is known to induce an inward rectifier potassium current (I<sub>KACH</sub>). This action of ACh is mediated by M<sub>2</sub> receptors coupled to I<sub>KACH</sub> via G<sub>i</sub> protein (Yatani et al., 1988; Takano and Noma, 1997). Recent

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**ABBREVIATIONS:** mAChR, muscarinic acetylcholine receptor; ACh, acetylcholine; TMA, tetramethylammonium; I<sub>KTMA</sub>, tetramethylammonium-induced K<sup>+</sup> currents; PTX, pertussis toxin; 4AP, 4-aminopyridine; I<sub>K4AP</sub>, 4-aminopyridine-induced K<sup>+</sup> currents; PZ, pirenzepine; Meth, methoctramine; 4-DAMP, 4-diphenylacetoxy-N-methylpiperidine methiodide; *p*-F-HHSiD, hexahydro-sila-difenidol hydrochloride, *p*-fluoro analog; Trop, tropicamide; [<sup>3</sup>H]NMS, [methyl-<sup>3</sup>H]N-scopolamine methyl chloride; *p*K<sub>i</sub>, equilibrium dissociation constant; RT, reverse transcription; PCR, polymerase chain reaction; I<sub>Kr</sub>, rapid component of delayed rectifier K<sup>+</sup> current; I<sub>Ks</sub>, slow component of delayed rectifier K<sup>+</sup> current.

studies have provided data showing the expression of other subtypes in cardiac tissues as well. For example, mRNAs coding for  $M_2$ ,  $M_3$ , and  $M_4$  isoforms have been shown to express in the chick heart (Tietje et al., 1990; Tietje and Nathanson, 1991; Gadbut and Galper, 1994; McKinnon and Nathanson, 1995). Sharma et al. (1996) identified and characterized the  $M_1$  receptors in the rat hearts. Functional data suggesting the presence of novel subtypes of mAChRs in cardiomyocytes have also been reported. For example, Fermini and Nattel (1994) first described a novel delayed rectifier-like  $K^+$  current activated by choline via the stimulation of mAChRs in canine atrial myocytes. Subsequently, Navarro-Polanco and Sánchez-Chapula (1997) demonstrated that 4-aminopyridine (4AP), a  $K^+$  channel blocker, was also able to activate a similar  $K^+$  current in cat atrial cells, an effect requiring the stimulation of mAChRs. Because these currents possess biophysical properties distinct from  $I_{K_{ACh}}$ , novel subtypes of mAChRs other than  $M_2$  were proposed by the authors as a mechanism underlying the activation of these channels. On the other hand, tetramethylammonium (TMA), a frequently used substance for replacing sodium ions in patch-clamp studies, has been demonstrated to be able to slow heart rate and weaken contractility via stimulation of mAChR in rat hearts (Kennedy et al., 1995). Workers from our laboratory and others have recently suggested that TMA, like 4AP, is also able to induce  $K^+$  currents by interacting with mAChRs (Navarro-Polanco and Sánchez-Chapula, 1998). However, it remains unclear which subtype of mAChRs mediates the actions of TMA.

Taken together, it is quite conceivable that other mAChR subtypes in addition to  $M_2$  also may exist in the heart. To date, however, little evidence has been submitted concerning the character and functionality of these subtypes in the heart. The goal of the present study was to explore mAChR subtypes in the canine heart using TMA and 4AP as pharmacological probes.

## Materials and Methods

**Cell Isolation.** Single canine atrial myocytes were isolated according to previously described techniques (Yue et al., 1997). The right atrium from adult mongrel dogs (20–26 kg) of either sex was quickly dissected and mounted via the right coronary artery to a Langendorff perfusion system. The preparation was perfused with  $Ca^{2+}$ -containing Tyrode's solution at 37°C until the effluent was clear of blood and then switched to  $Ca^{2+}$ -free Tyrode's solution for 20 min at a constant rate of 12 ml/min, followed by perfusion with the same solution containing collagenase (110 U/ml CLS II collagenase; Worthington Biochemical, Freehold, NJ) and 0.1% BSA (Sigma Chemical Co., St. Louis, MO). The dispersed cells were stored in KB medium at 4°C for later electrophysiological experiments.

**Solutions and Drugs.** The Tyrode's solution for cell isolation and whole-cell patch-clamp recording was composed of 136 mM NaCl, 5.4 mM KCl, 1 mM  $MgCl_2$ , 0.33 mM  $NaH_2PO_4$ , 5 mM HEPES, 10 mM glucose, and 1 mM  $CaCl_2$ , pH adjusted to 7.4 with NaOH. The KB medium for cell storage contained 20 mM KCl, 10 mM  $KH_2PO_4$ , 25 mM glucose, 70 mM potassium glutamate, 10 mM  $\beta$ -hydroxybutyric acid, 20 mM taurine, 10 mM EGTA, 0.1% albumin, and 40 mM mannitol, pH adjusted to 7.4 with KOH. The pipette solution contained 0.1 mM GTP, 110 mM potassium aspartate, 20 mM KCl, 1 mM  $MgCl_2$ , 5 mM Mg-ATP, 10 mM HEPES, 10 mM EGTA, and 5 mM phosphocreatine, pH adjusted to 7.3 with KOH. Contamination by sodium current was prevented by holding the cells at  $-50$  mV. Potential contamination by other currents was minimized by includ-

ing the following compounds in the bath solution: 1  $\mu$ M dofetilide (to inhibit  $I_{K_r}$ ), 20  $\mu$ M 293B (to block  $I_{K_A}$ ), 10  $\mu$ M glyburide (to prevent ATP-sensitive  $K^+$  current), and 200  $\mu$ M  $Cd^{2+}$  (to suppress calcium current). All chemicals were purchased from Sigma Chemical Co., except for 293B, which was a kind gift from Hoechst Pharmaceuticals (Frankfurt, Germany).

**Patch-Clamp Techniques.** Patch-clamp recording techniques used in this study have been described in detail elsewhere (Wang et al., 1994). Ionic currents were recorded with the whole-cell voltage-clamp methods using an Axopatch 200B amplifier (Axon Instruments, Burlingame, CA). Borosilicate glass electrodes (1 mm A) had tip resistances of 1 to 3 M $\Omega$  when filled with pipette solution. Junction potentials were zeroed before formation of the membrane-pipette seal in Tyrode's solution. Mean seal resistance averaged  $15 \pm 1$  G $\Omega$ . Several minutes after seal formation, the membrane was ruptured by gentle suction to establish the whole-cell configuration. The capacitance and series resistance ( $R_s$ ) was electrically compensated to minimize the duration of the capacitive surge on the current recording and the voltage drop across the clamped cell membrane.  $R_s$  along the clamp circuit was estimated by dividing the time constant obtained by fitting the decay of the capacitive transient by the calculated membrane capacitance (the time integral of the capacitive response to a 5-mV hyperpolarizing step from a holding potential of 0 mV divided by the voltage drop). Before  $R_s$  compensation, the decay of the capacitive surge was a single exponential function of time with a time constant of  $412 \pm 21$  ms (cell capacitance,  $82 \pm 5$  pF;  $n = 54$  cells). Precompensation  $R_s$  values averaged  $5.0 \pm 0.4$  M $\Omega$ . After compensation, the time constant was reduced to  $111 \pm 4$  ms (cell capacitance,  $71 \pm 4$  pF), and  $R_s$  was reduced to  $1.4 \pm 0.1$  M $\Omega$ . Currents recorded during the present study rarely exceeded 2.0 nA. The mean maximum voltage drop across the  $R_s$  was thus in the range of 3 mV. Cells with changing leak current (indicated by  $>10$  pA changes in holding current at  $-50$  mV) were rejected. Experiments were conducted at  $36 \pm 1^\circ$ C.

**Membrane Receptor-Binding Assay.** Fresh atrial tissues dissected from canine hearts were minced and washed with ice-cold PBS buffer. The tissues were then homogenized with a Polytron in 15 ml of ice-cold lysis buffer containing 5 mM Tris-HCl and 2 mM EDTA, pH 7.4, plus a protease inhibitor cocktail consisting of 5  $\mu$ g/ml phenylmethylsulfonyl fluoride, 10  $\mu$ g/ml benzamidine, and 5  $\mu$ g/ml soybean trypsin inhibitor. The homogenate was centrifuged at 500g for 15 min at 4°C. The pellets were then homogenized as before and spun again, and the supernatants were pooled. The supernatants were centrifuged at 45,000g for 15 min, and the pellets (membrane fractions) were washed twice in the same buffer. The membrane fractions were resuspended in a buffer containing 75 mM Tris-HCl, pH 7.4, 12.5 mM  $MgCl_2$ , and 5 mM EDTA. The protein content was determined with a Bio-Rad Protein Assay kit (Bio-Rad, Mississauga, ON) using BSA as the standard.

Saturation-binding assays were performed using eight concentrations of [*methyl*- $^3$ H]*N*-scopolamine methyl chloride ( $[^3$ H]NMS) ranging from 2 to 2500 pM. Nonspecific binding was measured in the presence of 1  $\mu$ M atropine. Experiments were carried out in triplicate for each experiment with total of four individual preparations. Incubations (90 min at room temperature) were terminated by rapid filtration using GF/B filters (Xymotech, Montreal, PQ), and radioactivity was counted with an LS6500 Scintillation Counter (Beckman, Fullerton, CA) with average efficiency of 58%.

Competition-binding assays were carried out as follows. Homogenates were incubated with 400 pM [ $^3$ H]NMS with pirenzepine (PZ; 0.3 nM to 100  $\mu$ M), methoctramine (Meth; 0.3 nM to 100  $\mu$ M), 4-diphenylacetoxy-*N*-methylpiperidine methiodide (4-DAMP; 0.1 nM to 100  $\mu$ M), or tropicamide (Trop; 1 nM to 100  $\mu$ M), TMA (30 nM to 1 mM), and 4AP (30 nM to 1 mM), respectively. Fixed amounts of membrane protein (100  $\mu$ g) were used for each sample in the binding study. Six individual experiments were performed with each determination performed in duplicate for each compound. Chemicals and

reagents for the binding study were purchased from Research Biochemicals International (Natick, MA)

**Reverse Transcription-Polymerase Chain Reaction.** RNA isolation and reverse transcription-polymerase chain reaction (RT-PCR) methods were the same as previously described (Wang et al., 1996, 1997, 1998). The total RNA samples extracted from canine atrial tissues were incubated with DNase I (0.1 U/ $\mu$ l) at 37°C for 15 min, followed by phenol/chloroform extraction to remove the genomic DNA. Integrity of total RNA was evaluated by ethidium bromide staining in denaturing agarose gels. RT was carried out in a 20- $\mu$ l reaction mixture containing 1 $\times$  reaction buffer (10 mM Tris-HCl, pH 8.3, 50 mM KCl, 2.5 mM MgCl<sub>2</sub>), 1 mM concentrations of dNTPs (Boehringer Mannheim, Montreal, Quebec, Canada), 3.2  $\mu$ g of random primers p(dN)<sub>6</sub> (Boehringer Mannheim), 5 mM dithiothreitol, 50 U of RNase inhibitor (Gibco BRL, Ontario, Canada), and 200 U of Moloney murine leukemia virus reverse transcriptase (Gibco BRL). First-strand cDNAs were synthesized at 42°C for 60 min, and the remaining enzymes were inactivated by heating at 99°C for 5 min. First-strand cDNA (5  $\mu$ l) resulting from RT was used as a template for amplification in a 25- $\mu$ l reaction mixture. Reagents included in each reaction contained 10 mM Tris-HCl, pH 8.3, 50 mM KCl, 1.5 mM MgCl<sub>2</sub>, 1 mM concentrations of dNTPs, 0.5  $\mu$ M concentration of each gene-specific primer, and 2.5 U of *Taq* polymerase (Gibco BRL). Reactions were hot-started at 94°C and continued for 3 min of initial melting. The cycling profiles were 30 s of denaturing at 94°C, 30 s of annealing at 50°C, and 40 s of extension at 72°C, for 30 cycles, followed by a final extension step of 5 min at 72°C.

The primers used in this study were for m<sub>1</sub> isoform, gggtgcagcagcagcagc (sense) and ctctttccacggggcttctg (antisense); for m<sub>2</sub> isoform, aagaaggacaagaaggagcc (sense) and ctttggatggcccagg (antisense); for m<sub>3</sub> isoform, tgcaggccagagaagc (sense) and ccttttcgcttagtgatctg (antisense); and for m<sub>4</sub> isoform, cccgaaggagaagaaagc (sense) and agtggtggcctctgtgg (antisense). They were synthesized by Gibco BRL. Because the canine cDNA sequences for mAChRs were not available, cloning of fragments was required. For this purpose, degenerate primers for varying isoforms of mAChRs were designed based on published sequences, and unique oligonucleotide sequences were chosen from cDNA regions bearing minimal homology among the different mAChR subtypes and minimal heterogeneity among the species for a given subtype. RT-PCR was used first to obtain the cDNA fragments from RNA extracts of canine right atrium. PCR products were size-separated on 1.5% agarose gels. Bands of desired size were purified with the Glassmax DNA Isolation Spin Cartridge System (Gibco BRL). The purified DNA fragments were subcloned into pGEM-T easy vector (Promega, Madison, WI). Sequences of all constructs were analyzed with Sequenase V2.0 (Amersham Life Science, Cleveland, OH). Gene-specific primer pairs were designed

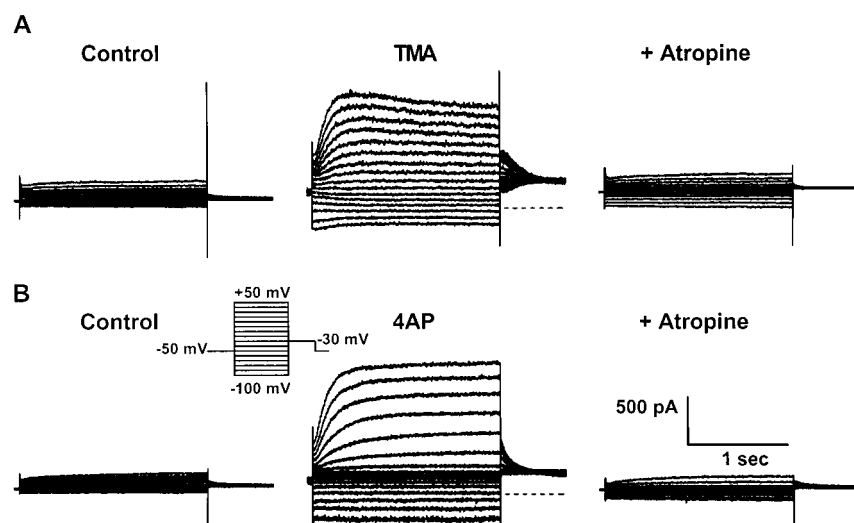
based on our cloned canine cDNA sequences. No sequences matching m<sub>1</sub> receptor cDNA could be obtained from dog hearts. Therefore, primers used for m<sub>1</sub> were designed from human sequence. All primers were designed to avoid significant secondary and complementary structures with about 50 to 60% G-C contents. Specificity of the primer pairs was verified by comparison with entire GenBank database using BLAST and FASTA and by amplification of a PCR product of the predicted size visible as a discrete single band with ethidium bromide staining on agarose gel.

To ensure that the RNA samples used in our experiments were free of contamination with RNA from sources other than cardiac tissue such as neurons or vascular cells, additional experiments were conducted and the detailed description is provided in *Results*.

**Data Analysis.** Group data are expressed as mean  $\pm$  S.E. Statistical comparisons were performed on raw data with Student's *t* test, with a two-tailed *p* < .05 taken to indicate a statistically significant difference. Binding data were analyzed using curve-fitting functions of the Prism software (GraphPAD Software, La Jolla, CA). To ensure validity and accuracy of displacement-binding studies, linear regression was performed on the percent bound versus the ratio of bound over free ligand and only data with a regression coefficient of 0.9 were accepted for analyses. The *F* test was used to compare fits for the competition-binding data, and the best fit (one-site binding versus two-site binding) was determined by the probability value for the *F* test and by the change in the residual sum of squares for the two different fits. One- and two-site models were tested for all data sets, and the model yielding the least residual sum of squares was taken to describe the data.

## Results

As illustrated in Fig. 1A, the currents conducting either outward or inward currents in response to depolarizing or hyperpolarizing voltage steps in the presence of TMA were consistently observed (middle and were not seen before exposure of the cells to TMA (left). TMA-induced outward currents were characteristic of delayed rectifier K<sup>+</sup> currents with slow time-dependent activation to a peak during depolarization and deactivation tail currents on repolarization to -30 mV. The effects of TMA were concentration-dependent (1  $\mu$ M to 10 mM), with detectable induction of the currents at a concentration of as low as 1  $\mu$ M and significant (*p* < .05, compared with baseline recordings) induction of the currents at 10  $\mu$ M. Maximal level of the current induction was reached at a concentration of 0.5 mM. Hence, 0.5 mM TMA was used in the remainder of the experiments.



**Fig. 1.** Activation of novel K<sup>+</sup> currents by TMA and 4AP via simulation of mAChRs in canine atrial myocytes. Currents were elicited by 2-s pulses to potentials ranging from -100 to +50 mV with 10-mV increment followed by a 1-s step to -30 mV. Voltage steps were delivered from a holding potential (HP) of -50 mV at an interspike interval of 5 s. A, raw current traces obtained from a representative myocyte, showing the baseline recordings before TMA application (left), currents activated in the presence of TMA (0.5 mM, *I*<sub>KTMA</sub>, middle), and elimination of the currents by atropine (1  $\mu$ M) applied to the bath containing TMA (right). B, typical examples of 4AP-induced currents recorded from a representative cell. Left, the baseline recordings before 4AP application, currents induced in the presence of 4AP (1 mM, *I*<sub>K4AP</sub>, middle), and abolishment of the currents by atropine (1  $\mu$ M) applied to the solution containing 4AP (right). Dashed line indicates a zero current level.



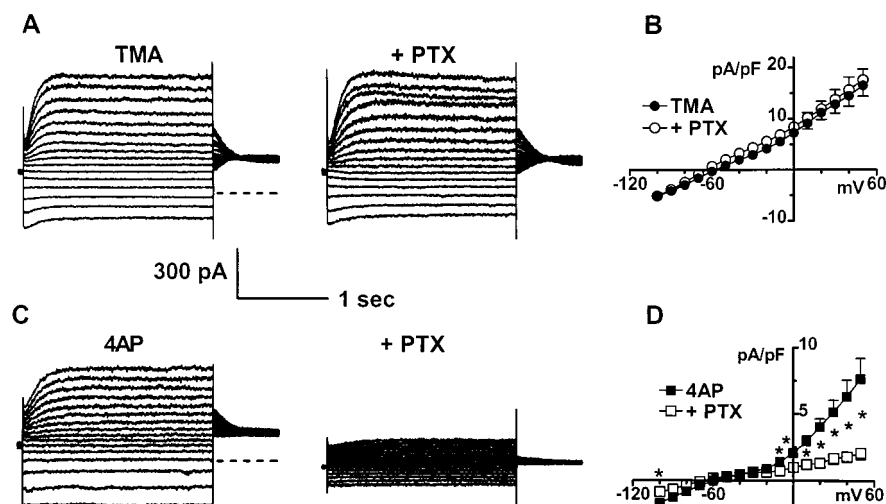
Similar to TMA, bath application of 4AP (1 mM) also consistently led to activation of voltage-dependent currents similar to TMA-induced currents with conductance in both outward (depolarizing pulses) and inward (hyperpolarizing pulses) directions. Representative current traces are shown in Fig. 1B. The size of 4AP-induced currents also depended on 4AP concentration (50  $\mu$ M to 2 mM), with maximal effects occurring at a concentration of 1 mM. Thus, 1 mM 4AP was used to study the 4AP-induced currents in the remainder of our experiments. There was no apparent run-down of TMA- or 4AP-induced currents under our experimental conditions with stable whole-cell recordings up to 2 h. In both cases, the current induction was seen right after (within 3 min) the wash-in of the drugs into the bath.

Both currents reversed at around the  $K^+$  equilibrium potential ( $-80$  mV) according to Nernst equation under our experimental conditions ( $-73.4 \pm 5.6$  mV for TMA-induced currents,  $n = 3$ ; and  $-70.2 \pm 7.2$  mV for 4AP-induced currents,  $n = 3$ ). Furthermore, the reversal potentials of both TMA- and 4AP-induced currents, as determined by the tail currents elicited at various potentials ranging from  $-120$  to  $-30$  mV, became less negative with elevating concentrations of external  $K^+$  (isotonic replacement of  $Na^+$  with  $K^+$  from 5.4 to 10.8 and 130 mM, respectively). The calculated slope factor from linear regression of reversal potentials (as a function  $[K^+]_o$ ) were  $51.3 \pm 3.3$  and  $50.2 \pm 5.2$  mV/decade for TMA- and 4AP-induced currents, respectively, suggesting that both currents were predominantly carried by  $K^+$ .

To investigate whether the currents were resulted from TMA/4AP modulation of known  $K^+$  currents, we evaluated the effects of varying  $K^+$  channel blockers on TMA- and 4AP-induced  $K^+$  currents. These blockers included glyburide (10  $\mu$ M for  $I_{KATP}$ ), dofetilide (1  $\mu$ M for  $I_{Kr}$ ), indapamide (100  $\mu$ M for  $I_{Ks}$ ), 293B (20  $\mu$ M for  $I_{Ks}$ ), and  $Cd^{2+}$  (200 mM for  $I_{Ca}$ ). We failed to observe any significant alterations of TMA- and 4AP-induced currents in the presence of any one of these compounds, suggesting that these two currents are distinct from the known potassium currents previously found in the heart. We therefore included glyburide, dofetilide, 293B, and  $Cd^{2+}$  in the perfusate for the remainder of our experiments. For the sake of convenience, we used abbreviated labels  $I_{KTMA}$  and  $I_{K4AP}$  to describe the TMA- and 4AP-induced  $K^+$  currents, respectively, in the remainder of the article.

The above data indicated that  $I_{KTMA}$  and  $I_{K4AP}$  probably represent novel currents similar to recently described  $K^+$  currents in dog atrium (Fermini and Nattel, 1994) and cat atrium (Navarro-Polanco and Sánchez-Chapula, 1997), respectively, for which the activation of mAChRs was required. Indeed, atropine (1  $\mu$ M), a nonselective mAChR antagonist, quickly abolished  $I_{KTMA}$  and  $I_{K4AP}$ . The results are displayed in Fig. 1 (right). Similar results were observed in total of 11 cells for  $I_{KTMA}$  and 10 cells for  $I_{K4AP}$ .

The atropine data indicate that stimulation of mAChRs is required for the activation of  $I_{KTMA}$  and  $I_{K4AP}$ . It is known, however, that an  $M_2$  receptor is the only mAChR subtype functionally identified in cardiac cells (Yatani et al., 1988; Takano and Noma, 1997) and that stimulation of an  $M_2$  receptor by ACh or carbachol induces an inward rectifier  $K^+$  current,  $I_{KACH}$ . If it is true that only  $M_2$  receptors exist in the heart, then why would TMA- or 4AP-induced current demonstrate delayed rectifying properties distinct from  $I_{KACH}$ ? One of the possible explanations for this apparent contradiction is that there are other subtypes of mAChRs in addition to  $M_2$  receptors in canine atrial myocytes, which mediate the activation of  $I_{KTMA}$  and  $I_{K4AP}$ . We therefore explored the possible role of mAChR subtypes in canine atrial myocytes. We first carried out experiments to evaluate the effects of pertussis toxin (PTX) on  $I_{KTMA}$  and  $I_{K4AP}$ . PTX was applied intracellularly through the dialysis of pipette solution containing 2  $\mu$ g/ml PTX. Currents were recorded immediately after membrane rupture and taken as baseline values. Then, same recordings were repeated every 5 min, and currents recorded 20 min after membrane rupture were used for data analysis representing steady-state effects of PTX because 10 min was sufficient to allow the complete dialysis of PTX into the cellular plasma under our experimental conditions. No significant changes ( $p > .05$ , compared with baseline values,  $n = 5$  cells) in  $I_{KTMA}$  were observed in the presence of PTX (Fig. 2A). In sharp contrast,  $I_{K4AP}$  was nearly abolished ( $p < .05$ ,  $n = 5$ ) by PTX, as shown in Fig. 2C. For comparison, the effects of PTX on  $I_{KACH}$  were also determined, and similar results to those seen with  $I_{K4AP}$  were observed. As mentioned, neither  $I_{KTMA}$  nor  $I_{K4AP}$  demonstrated any rundown during the recording. This validates our PTX experiments described above. To further confirm the observations, the effects of extracellularly applied PTX on  $I_{KTMA}$  and  $I_{K4AP}$



**Fig. 2.** Effects of PTX, a  $G_i/G_o$  protein inhibitor, on  $I_{KTMA}$  and  $I_{K4AP}$  in canine atrial myocytes. PTX (2  $\mu$ g/ $\mu$ l) was applied intracellularly by dialysis through the pipette solution. A, analog data shows  $I_{KTMA}$  traces recorded immediately after membrane rupture (left) and 20 min after membrane rupture (middle) in the presence of TMA. Right, current density-voltage relation of  $I_{KTMA}$  averaged from five cells. B, current density-voltage relationship of  $I_{KTMA}$  before and after PTX treatment (averaged data from a total of five cells tested). C, effects of PTX on  $I_{K4AP}$ . Shown is a family of current traces recorded in the presence of 4AP immediately after membrane rupture (left) and 20 min after membrane rupture (middle). D, current density-voltage relationship of  $I_{K4AP}$  before and after PTX treatment ( $n = 5$ ). Dashed line indicates a zero current level. \* $p < .05$  comparison between before and after PTX.

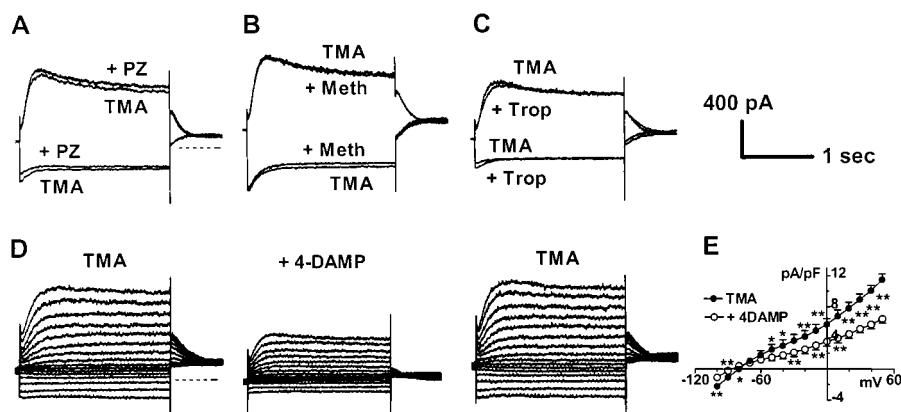
were also assessed. In these experiments, cells were divided into two groups: the control group and the PTX group. The cells of the PTX group were preincubated with a final concentration of 5  $\mu\text{g/ml}$  PTX in KB solution at 36°C for 90 min. The control cells were subjected to the same procedure but without PTX in the solution. Then, patch-clamp recording was carried out to study the inducibility of  $I_{\text{KTMA}}$  or  $I_{\text{K4AP}}$  in cells from the PTX group compared with cells from the control group. Consistent with the results from intracellular application of PTX, pretreatment with PTX did not alter TMA (0.5 mM) induction of  $I_{\text{KTMA}}$  compared with the control cells. For example, the  $I_{\text{KTMA}}$  density, measured at +50 mV, was  $9.8 \pm 1.3$  pA/pF for the control group ( $n = 8$ ) and  $9.6 \pm 0.8$  pA/pF for the PTX group ( $n = 7$ ,  $p > .05$ ). In contrast, the ability of 4AP to induce  $I_{\text{K4AP}}$  was completely paralyzed by PTX. The current density of  $I_{\text{K4AP}}$  was  $6.3 \pm 0.6$  pA/pF ( $n = 3$ ) for the control cells, and this value was decreased to  $0.1 \pm 0.0$  pA/pF ( $n = 3$ ,  $p < .01$ ) with the PTX-pretreated cells.

The results of the PTX experiments suggest that  $I_{\text{KTMA}}$  is probably mediated by an M<sub>1</sub>, M<sub>3</sub>, or M<sub>5</sub> receptor subtype because these subtypes are characterized biochemically by coupling to inositol phosphate production via a PTX-insensitive G protein (Peralta et al., 1988), whereas  $I_{\text{K4AP}}$  is likely mediated by M<sub>2</sub> or M<sub>4</sub> subtype because the transduction machinery for these subtypes has been shown to be PTX sensitive (Peralta et al., 1988). To further pinpoint the mAChR subtype specificity of  $I_{\text{KTMA}}$  and  $I_{\text{K4AP}}$ , we explored the relationship between the currents and the receptor subtypes by using pharmacological probes selective to different mAChR subtypes. No appreciable alterations in  $I_{\text{KTMA}}$  were found after the application with PZ (an M<sub>1</sub>-selective antagonist; 10 nM) (Watson et al., 1983; van Zwieten and Doods, 1995), Meth (an M<sub>2</sub>-selective antagonist; 20 nM) (Michel and Whiting, 1988; van Zwieten and Doods, 1995), or Trop (an M<sub>4</sub>-selective antagonist; 200 nM) (Lazareno et al. 1990; Lazareno and Birdsall, 1993). No attempt was made to study the effects of higher concentrations of these compounds because subtype specificity is lost at higher concentrations. 4-DAMP, an M<sub>3</sub>-selective antagonist (Barlow and Shepherd, 1986; Michel et al., 1989; Araujo et al., 1991; van Zwieten and Doods, 1995), produced concentration-dependent suppres-

sion of  $I_{\text{KTMA}}$ . As shown in Fig. 3, the currents were reduced by approximately 50% at a concentration as low as 2 nM for both depolarization- and hyperpolarization-induced currents (Table 1), and complete inhibition was achieved with 10 nM 4-DAMP. In the case of  $I_{\text{K4AP}}$ , as shown in Fig. 4, only the M<sub>4</sub> inhibitor Trop caused significant depression of the current amplitude without changing the kinetics. The current density at +50 mV was reduced  $46 \pm 3\%$  by 200 nM,  $68 \pm 5\%$  by 1  $\mu\text{M}$ , and  $93 \pm 8\%$  by 10  $\mu\text{M}$  Trop ( $n = 5$  for each). Other compounds, like PZ (10 nM), Meth (20 nM), and 4-DAMP (2 nM), failed to alter  $I_{\text{K4AP}}$ .

Our data suggest that  $I_{\text{KTMA}}$  might be mediated by M<sub>3</sub> receptors and that  $I_{\text{K4AP}}$  might be mediated by M<sub>4</sub> receptors. To acquire more conclusive evidence, we assessed the effects of 4-DAMP and Trop on the concentration-dependent responses to TMA (1  $\mu\text{M}$  to 2 mM) and 4AP (10  $\mu\text{M}$  to 2 mM), respectively. The current density obtained at +50 mV was used to determine the drug effects. As shown in Fig. 5A, 4-DAMP (2 nM) caused a parallel shift of the concentration-response curve of  $I_{\text{KTMA}}$  to the right. The  $\text{EC}_{50}$  value of TMA was 129  $\mu\text{M}$  with a Hill coefficient of 2.3 in the absence of 4-DAMP and 447  $\mu\text{M}$  ( $P < .05$ ,  $n = 3$ ) with a Hill coefficient of 1.7 in the presence of 4-DAMP. Similar effects of Trop on  $I_{\text{K4AP}}$  were observed (Fig. 5B). The  $\text{EC}_{50}$  value of 4AP for  $I_{\text{K4AP}}$  induction was 181  $\mu\text{M}$  for 4AP alone versus 391  $\mu\text{M}$  ( $P < .05$ ,  $n = 3$ ) for 4AP in the presence of Trop (200 nM). The calculated Hill coefficient was 1.3 for both with and without Trop.

We also tested the ability of two other M<sub>3</sub>-selective inhibitors: 4-DAMP mustard (an irreversible M<sub>3</sub> antagonist; 4 nM,  $n = 5$ ) (Ehlert and Griffin, 1988) and hexahydro-siladifenidol hydrochloride, *p*-fluoro analog (*p*-F-HHSiD; 20 nM,  $n = 5$ ) (Lambrecht et al., 1988), to suppress  $I_{\text{KTMA}}$ . We observed  $49.4 \pm 3.3\%$  reduction in current at +50 mV by 4-DAMP mustard ( $p < .01$ ,  $n = 3$ ) and  $56 \pm 8\%$  reduction in current by *p*-F-HHSiD ( $p < .05$ ,  $n = 3$ ). It is generally agreed that the use of an irreversible receptor inactivation plus concurrent receptor protection can enhance selective alkylation (Ehlert and Griffin, 1988; Eglen et al., 1994). In our experiments, we first incubated the cells with antagonists for M<sub>1</sub>/M<sub>2</sub>/M<sub>4</sub> subtypes to protect these receptors, if any, and



**Fig. 3.** Effects of mAChR subtype-selective antagonists on  $I_{\text{KTMA}}$  in canine atrial myocytes. Currents were elicited in the presence of 0.5 mM TMA, with voltage protocols same as described in Fig. 1. A, effects of M<sub>1</sub>-selective antagonist PZ (10 nM). Raw current traces before and after PZ are superimposed, and only currents recorded at +50 mV and -100 mV are shown for the sake of clarity. B, effects of M<sub>2</sub>-selective antagonist Meth (20 nM). C, effects of M<sub>4</sub>-selective antagonist Trop (200 nM). D, representative experiments showing the effects of M<sub>3</sub>-selective antagonist 4-DAMP on TMA-induced K<sup>+</sup> currents. Left, currents before wash-in of 4-DAMP. Middle, currents in the presence of 4-DAMP (2 nM). Right, currents after washout of 4-DAMP. E, current density-voltage relationship of baseline current and after exposure of cells to 4-DAMP at a concentration of 2 nM. Dashed line indicates a zero current level. \* $p < .05$  and \*\* $p < .01$ , comparison between before and after 4-DAMP ( $n = 6$ ).

then applied alkylating agent 4-DAMP mustard to inactivate the unprotected subtype (i.e.,  $M_3$  receptors). The same degree of  $I_{KTMA}$  suppression by 4-DAMP mustard was seen with and without protection, strongly suggesting the role of  $M_3$  subtype in canine myocytes.

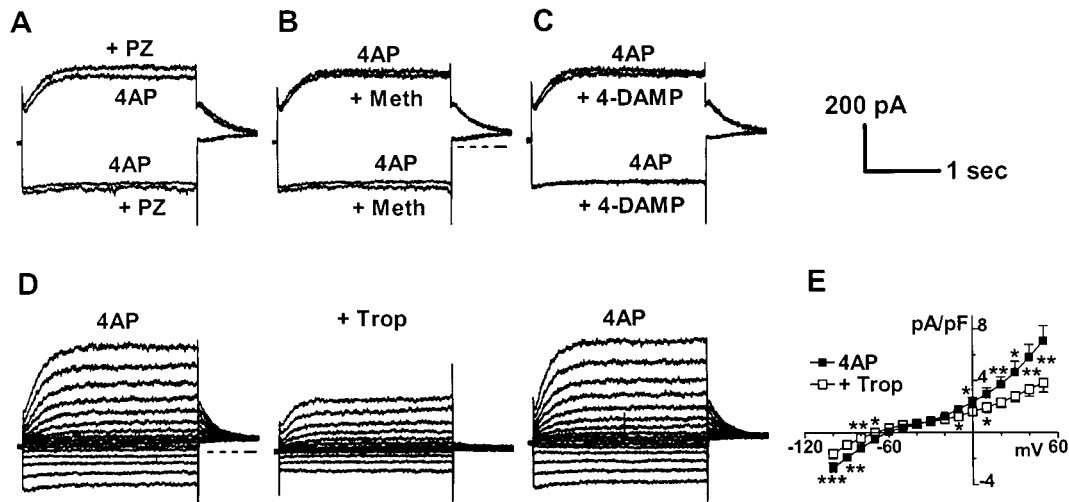
It is known that  $M_2$  subtype mediates ACh-induced inward rectifier  $K^+$  current ( $I_{KACH}$ ) in cardiac cells. Consistent with this notion, none of the antagonists except for Meth, an  $M_2$ -selective inhibitor, had any detectable effects on  $I_{KACH}$  at concentrations tested in our experiments (Fig. 6). By comparison, Meth at 20 nM substantially diminished  $I_{KACH}$  (Table 1). Meth (20 nM) did not alter the concentration-response curves of  $I_{KTMA}$  and  $I_{K4AP}$  (Fig. 5), suggesting a minimal

contribution of  $M_2$  to these currents. An immediate question that one might ask is whether ACh, as a nonselective mAChR agonist, could also activate  $I_{KTMA}$  and  $I_{K4AP}$ . As shown in Fig. 6D, when Meth was added to the ACh-containing solution,  $I_{KACH}$  was reduced due to inhibition of  $M_2$  receptors. Meanwhile, there was an increase in the outward currents. The outward portion of  $I_{KACH}$  without Meth had strong inward rectification, but in the presence of Meth, the outward currents displayed outward rectification, and this is clearly seen with the current-voltage curves shown in Fig. 6E. To determine whether these outward currents were the results of activation of  $M_3/M_4$  receptors by ACh unmasked when  $M_2$  receptors were largely inhibited, we evaluated the effects of

TABLE 1  
Effects of mAChR subtype-selective antagonists on  $I_{KTMA}$ ,  $I_{K4AP}$ , and  $I_{KACH}$

		Pirenzepine, 10 nM	Methoctramine, 20 nM	4-DAMP, 2 nM	Tropicamide, 200 nM
		pA/pF			
$I_{KTMA}$ +50 mV	Control	13.9 ± 2.1	12.7 ± 2.7	11.5 ± 0.6	9.0 ± 1.1
	Drug	14.0 ± 2.0	12.2 ± 2.7	6.4 ± 0.4**	9.1 ± 1.2
	% change	1.0 ± 3.5	-4.0 ± 0.6	-42.8 ± 6.1	0.5 ± 1.7
	-100 mV				
	Control	4.2 ± 0.9	4.1 ± 1.0	2.2 ± 0.3	3.5 ± 0.8
	Drug	4.3 ± 0.5	4.2 ± 0.9	1.1 ± 0.2**	3.8 ± 1.0
$I_{K4AP}$ +50 mV	Control	5.8 ± 8.2 (n = 4)	3.1 ± 4.0 (n = 5)	-50.9 ± 5.1 (n = 6)	5.2 ± 7.4 (n = 4)
	% change				
	-100 mV				
	Control	6.8 ± 1.5	7.8 ± 0.9	8.7 ± 0.2	7.1 ± 1.0
	Drug	6.9 ± 1.6	8.0 ± 0.8	9.0 ± 0.5	3.8 ± 0.5**
	% change	1.5 ± 3.8	-3.7 ± 0.9	7.2 ± 3.4	-46.4 ± 3.1
$I_{KACH}$ +50 mV	Control	3.8 ± 1.1	3.4 ± 1.0	3.8 ± 1.5	2.7 ± 0.3
	Drug	4.1 ± 1.2	3.8 ± 1.6	3.7 ± 1.3	1.6 ± 0.2***
	% change	5.9 ± 3.0 (n = 3)	12.9 ± 4.4 (n = 3)	-0.7 ± 5.1 (n = 4)	-41.8 ± 4.5 (n = 5)
	-100 mV				
	Control	7.0 ± 1.0	6.4 ± 0.6	5.9 ± 0.9	6.2 ± 0.5
	Drug	6.9 ± 1.2	9.4 ± 1.0**	5.7 ± 0.5	6.1 ± 0.3
$I_{KACH}$ -100 mV	% change	-0.2 ± 0.3	44.8 ± 4.0	-3.4 ± 0.5	-1.6 ± 1.0
	Control	11.2 ± 3.0	9.4 ± 2.1	10.4 ± 1.3	9.1 ± 1.8
	Drug	10.9 ± 4.1	14.8 ± 1.1	9.8 ± 3.5	8.9 ± 2.1
	% change	-1.0 ± 0.4 (n = 3)	-57.4 ± 4.7*** (n = 6)	-5.2 ± 1.2 (n = 3)	-2.2 ± 2.2 (n = 3)

\*  $p < .05$ , \*\*  $p < .01$ , \*\*\*  $p < .001$ . Student's  $t$  test, compared with control.



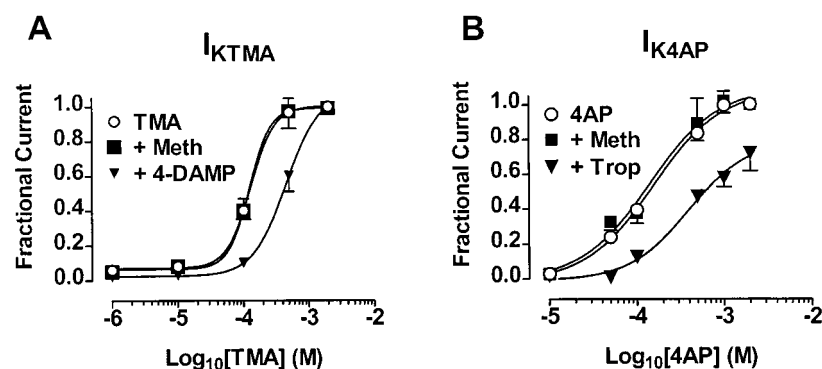
**Fig. 4.** Effects of mAChR subtype-selective antagonists on  $I_{K4AP}$  in canine atrial myocytes. Currents were elicited in the presence of 1 mM 4AP. A through C, effects of PZ (10 nM), Meth (20 nM), and 4-DAMP (2 nM) on  $I_{K4AP}$ , respectively. Raw current traces before and after the drugs are superimposed, and only currents recorded at +50 mV and -100 mV are shown for the sake of clarity. D, representative experiments showing the effects of  $M_4$ -selective antagonist Trop on  $I_{K4AP}$ . Left, currents before washout of Trop. Middle, currents in the presence of Trop (200 nM). Right, currents after washout of Trop. E, current density-voltage relationship of baseline current and after exposure of cells to Trop at a concentration of 200 nM. Dashed line indicates a zero current level. \* $p < .05$ , \*\* $p < .01$ , and \*\*\* $p < .01$ , comparison between before and after Trop.

4-DAMP or Trop on these outward currents. The results from representative experiments are illustrated in Fig. 7. Our data demonstrated that both 4-DAMP (2 nM) and Trop (200 nM) were able to depress the currents and the 4-DAMP- or Trop-sensitive currents, obtained by digital subtraction between the currents with and without 4-DAMP or Trop, had waveforms similar to  $I_{KTMA}$  or  $I_{K4AP}$ , suggesting that  $M_3$  and  $M_4$  receptors were involved in the activation of ACh-induced outward currents in the presence of Meth. The extents of block by 4-DAMP or Trop varied greatly, presumably depending on the relative amount of  $M_3$  or  $M_4$  receptors activated in each individual cell. It should be emphasized that the outward currents induced by ACh in the presence of Meth were observed only from 16 of 29 cells studied. To investigate whether the outward currents induced by ACh were actually due to nonspecific or some unknown effects of Meth, we applied Meth (20 nM) alone to the bath. Meth itself failed to induce any kinds of ion currents (Fig. 7C). Furthermore, when the cells were preincubated with 4-DAMP and Trop to prevent the stimulation of  $M_3$  and  $M_4$  receptors, ACh failed to induce the outward currents even after addition of Meth to the solution (Fig. 7D).

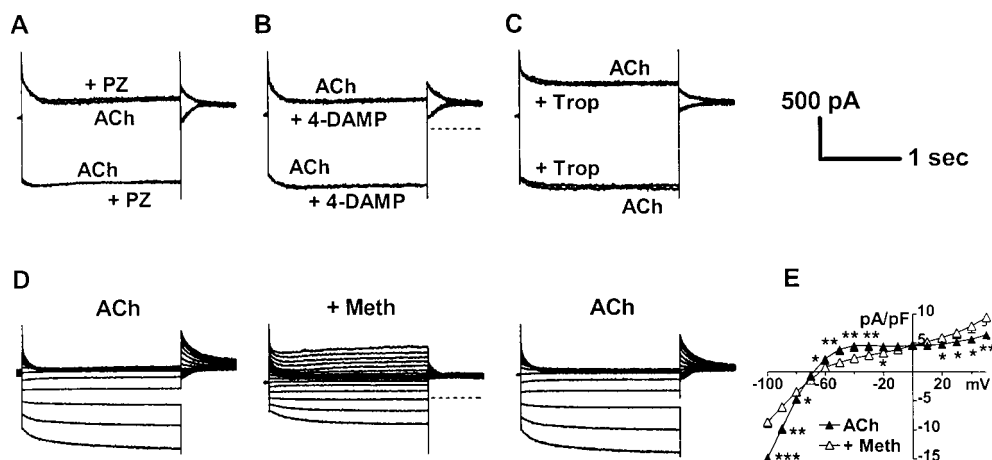
**Existence of  $M_3$  and  $M_4$  Receptors in Cardiac Cells Demonstrated by Binding Assay.** mAChR subtypes demonstrate characteristic affinities for the binding of different

muscarinic antagonists. To verify the presence of  $M_3$  and  $M_4$  receptor proteins in cardiac cells, we performed membrane receptor-binding assays. The results are summarized in Fig. 8 and Table 2. Saturation binding of [ $^3$ H]NMS to the membrane homogenates from canine atria yielded a maximum binding value or mAChR density of  $282 \pm 26$  fmol/mg protein and a dissociation constant  $K_d$  of  $223 \pm 24$  pM. Displacement binding of [ $^3$ H]NMS in the presence of PZ, Meth, 4-DAMP, or Trop was analyzed with a two-site binding model. The  $pK_i$  values from competition-binding experiments are listed in Table 2. Note that the  $K_d$  values for high-affinity bindings of various antagonists were all closely relevant to the concentrations at which these compounds produced half-maximal inhibition to the currents. For example, the  $K_d$  value of 4-DAMP was 2.4 nM, and 2 nM 4-DAMP suppressed about 50% of  $I_{KTMA}$ . In addition, the  $K_d$  value of Trop binding and the concentration of Trop that caused 50% reduction of  $I_{K4AP}$  were 230 and 200 nM, respectively.

If, as our functional study suggested, TMA and 4AP activated  $I_{KTMA}$  and  $I_{K4AP}$  via stimulation of  $M_3$  and  $M_4$  receptors, respectively, then these two compounds should also be able to displace [ $^3$ H]NMS binding in a competitive fashion. Indeed, the competition-binding assay performed with TMA and 4AP yielded displacement curves best described with a two-site binding model. The data are shown in Fig. 8C. The



**Fig. 5.** Effects of mAChR subtype-selective antagonists on the concentration dependence of TMA induction of  $I_{KTMA}$  (A) and 4AP-induced  $I_{K4AP}$  (B) in canine atrial myocytes. Symbols represent values averaged from three cells, and the lines represent the best fits to the Hill equation:  $B/B_{max} = 1/[1 + (EC_{50}/D)^n]$ , where  $B_{max}$  is the maximum level of currents inducible by agonists (e.g., 2 mM TMA or 2 mM 4AP),  $B$  is the current amplitude induced at a drug concentration  $D$ ,  $EC_{50}$  is the concentration of agonists that produces 50% maximum induction of the currents, and  $n$  is the Hill coefficient. Note that 4-DAMP (2 nM) or Trop (200 nM) markedly shifts the dose-response curve of  $I_{KTMA}$  or  $I_{K4AP}$  to the right, but Meth (20 nM) does not alter the concentration dependence of either of the currents.



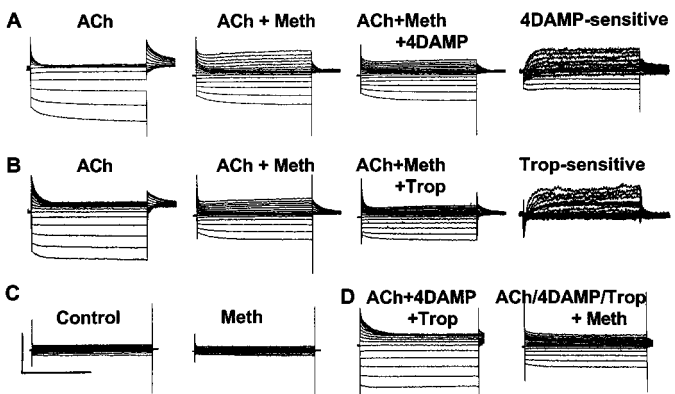
**Fig. 6.** Effects of mAChR subtype-selective antagonists on  $I_{KACh}$  in canine atrial myocytes. Currents were elicited in the presence of 1  $\mu$ M ACh, with the same voltage protocols as described in Fig. 1. A through C, effects of mAChR subtype-selective antagonists PZ (10 nM), 4-DAMP (2 nM), and Trop (200 nM) on  $I_{KACh}$ . Raw current traces before and after the drugs are superimposed, and only recordings obtained at +50 mV and -100 mV are shown. D, representative experiments showing the effects of  $M_2$ -selective antagonist Meth (20 nM) on  $I_{KACh}$ . Left, currents before wash-in of Meth. Middle, currents in the presence of Meth (20 nM). Right, currents after washout of Meth. E, current density-voltage relationship of baseline current and after exposure of cells to 20 nM Meth. Dashed line indicates a zero current level. \* $p < .05$ , \*\* $p < .01$ , and \*\*\* $p < .001$ , comparison before and after Meth.



$K_d$  values were 10.9  $\mu$ M and 2.5 mM for TMA and 2.5  $\mu$ M and 0.4 mM for 4AP. These values were both comparable to the concentrations at which these compounds produced induction of  $I_{KTMA}$  and  $I_{K4AP}$ , respectively.

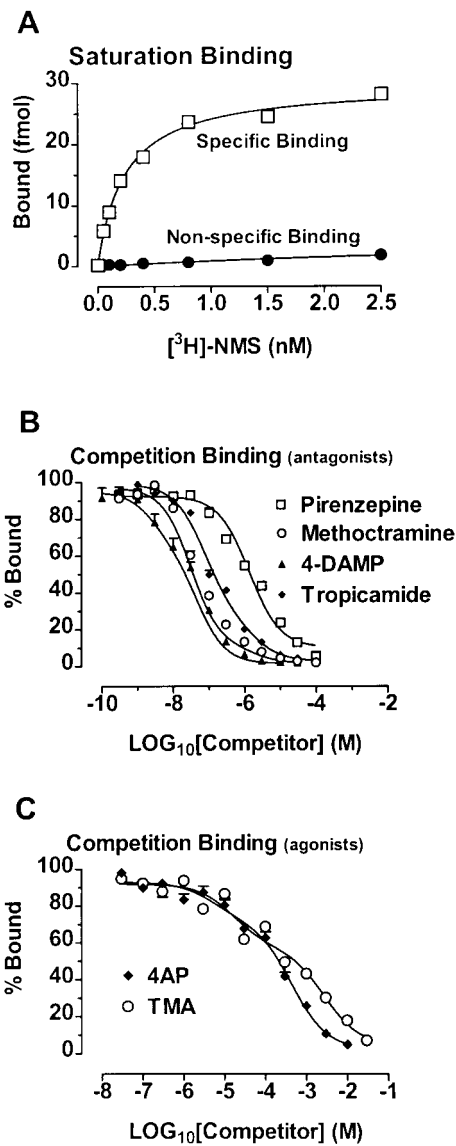
**Expression Profile of mRNA Coding for Various mAChR Isoforms.** To obtain further evidence for the presence of multiple subtypes of mAChRs in canine hearts, cloning of cDNA fragments and detection of mRNAs for  $m_1$ ,  $m_2$ ,  $m_3$ , and  $m_4$  four different isoforms of mAChRs were performed by RT-PCR amplification. We isolated cDNA fragments of 458 bp for  $m_2$ , 432 bp for  $m_3$ , and 258 bp for  $m_4$  isoforms, but we failed to obtain any sequences matching with  $m_1$  cDNA. These fragments share 91%, 81%, and 80% homology to the same regions of corresponding human  $m_2$ ,  $m_3$ , and  $m_4$  sequences in the amino acid level. These fragments represent parts of the third intracellular loop between transmembrane domains 5 and 6, which is thought to contain critical determinants of G protein-coupling specificity. A representative gel showing the PCR-amplified products representing mRNAs for  $m_2$ ,  $m_3$ , and  $m_4$  subtypes is shown in Fig. 9. The  $m_2$ ,  $m_3$ , and  $m_4$  isoforms were all expressed at significant levels, as indicated by the 297-bp band in lane 4 for  $m_2$ , the 216-bp band in lane 6 for  $m_3$ , and the 257-bp band in lane 8 for  $m_4$ .

Previous investigators have assayed mRNA for the neuronal acetylcholine  $\beta_4$  subunit to exclude neural contamination of cardiac mRNA measurements (Dixon and McKinnon, 1994). We used published sequences of the human nicotinic receptor  $\beta_4$  subunit to design primer pairs for RT-PCR. The  $\beta_4$  subunit mRNA was consistently detected in three samples from rat brain but was absent in dog atrial samples. Con-



**Fig. 7.** Effects of 4-DAMP and Trop on the outward currents induced by ACh in the presence of Meth. A, examples showing 4-DAMP (2 nM) blockade of ACh (1  $\mu$ M)-induced outward currents in the presence of Meth (20 nM) from a representative myocyte. The 4-DAMP-sensitive currents were obtained by digital subtraction of the currents with 4-DAMP from those before 4-DAMP. Similar results were obtained in another four cells. B, examples showing Trop (200 nM) blockade of ACh (1  $\mu$ M)-induced outward currents in the presence of Meth (20 nM) from a representative myocyte. The Trop-sensitive currents were obtained by digital subtraction of the currents in the presence of Trop from those before Trop. Similar data were obtained from a total of five cells. C, experiments demonstrating the failure of Meth (20 nM) alone to induce any currents. The same observations were obtained from all five cells studied. D, experiments showing the inability of ACh to induce the outward current when 4-DAMP (2 nM) and Trop (200 nM) were included in the solution. The addition of Meth still failed to activate the outward current seen in the absence of 4-DAMP and Trop but reduced  $I_{KACh}$ . Similar results were acquired from another three myocytes. Horizontal calibration, 1 s. Vertical calibration for all except for the 4-DAMP- and Trop-sensitive currents was 500 pA, and that for the 4-DAMP- and Trop-sensitive currents was 200 pA.

tamination by vascular tissue was excluded by RT-PCR amplification of the maxi-K channel ( $Ca^{2+}$ -activated  $K^+$  channel). Total RNA isolated from rat vascular smooth muscle, in which maxi-K channels carry a substantial current, was used



**Fig. 8.** mAChR binding with [ $^3$ H]NMS to homogenates of canine atrial tissues. A, saturation binding. Shown are the data averaged from four individual preparations carried out in triplicate for each experiment. Symbols are experimental data and lines represent the fits with a one-site binding model. B, displacement binding of [ $^3$ H]NMS with varying antagonists. C, displacement binding of [ $^3$ H]NMS with two agonists: TMA and 4AP. Each point represents the mean  $\pm$  S.E.M. from six experiments assayed in duplicate. Curves are best fits of the experimental data to a two-site binding model.

**TABLE 2**  
Displacement binding of [ $^3$ H]NMS by mAChR subtype-selective antagonists and agonists

	$K_{i1}$	$K_{i2}$
Pirenzepine	$6.9 \pm 0.3$	$5.8 \pm 0.2$
Methoctramine	$7.7 \pm 0.6$	$6.6 \pm 0.4$
4-DAMP	$9.1 \pm 0.2$	$8.1 \pm 0.1$
Tropicamide	$7.6 \pm 0.1$	$6.4 \pm 0.2$
TMA	$5.7 \pm 0.2$	$3.1 \pm 0.0$
4AP	$7.0 \pm 0.5$	$3.9 \pm 0.1$

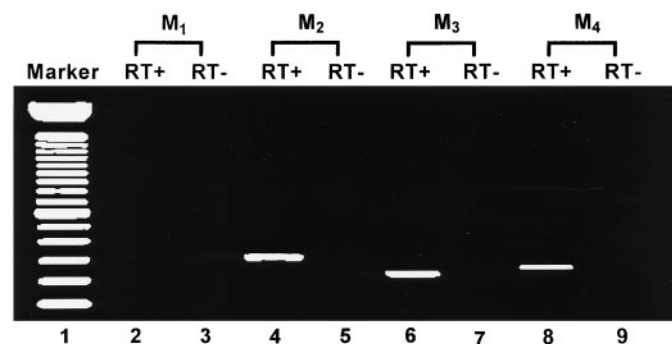


as a positive control. PCR primer pairs were designed based on published sequence. Although strong maxi-K channel signal was observed in vascular preparations, no corresponding band was found in dog tissue.

## Discussion

One major breakthrough in the field of the cholinergic nervous system is the discovery of multiple subclasses of muscarinic receptors, due to the development of pharmacological probes and molecular cloning techniques. Although M<sub>2</sub> receptors have long been believed to be the only functional mAChR subtype in cardiac cells, our data from functional study (patch-clamp experiments), receptor-binding assays, and mRNA expression detection demonstrate the heterogeneity of mAChRs and their physiological functions in canine atrial cells. There are two major and novel findings in the present study. First, multiple mAChR subtypes (M<sub>2</sub>, M<sub>3</sub>, and M<sub>4</sub>) coexist in canine cardiac cells. Our data represent, to our knowledge, the first to identify and characterize cardiac M<sub>3</sub> and M<sub>4</sub> receptor subtypes in the heart with both functional and binding data. Second, different subtypes of mAChRs are coupled to different K<sup>+</sup> channels in cardiac myocytes.

To date, at least four subtypes of mAChRs have been pharmacologically defined in living systems: M<sub>1</sub>, M<sub>2</sub>, M<sub>3</sub>, and M<sub>4</sub> (Eglen and Whiting, 1990; Mutschler et al., 1995; van Zwieten and Doods, 1995), and five different isoforms of mAChR cDNAs have also been molecularly identified, which are designated as m<sub>1</sub> through m<sub>5</sub> (Bonner et al., 1987; Peralta et al., 1987; Goyal, 1989). It has long been believed that M<sub>2</sub> is the only functional mAChR subtype in cardiac cells (Watson et al., 1983; Bonner et al., 1987; Mizushima et al., 1987; Peralta et al., 1987; Maeda et al., 1988). However, the existence and possible role of M<sub>1</sub> receptors in the rat hearts have been convincingly documented by Sharma et al. (1996). Recent studies with molecular cloning techniques also reported interesting data showing the abundant expression of mRNAs for m<sub>2</sub>, m<sub>3</sub>, and m<sub>4</sub> subtypes in the chick heart (Tietje et al., 1990; Tietje and Nathanson, 1991; Gadbut and Galper, 1994; McKinnon and Nathanson, 1995). In the present study, we provide three lines of evidence for the coexistence of multiple subtypes of mAChRs in cardiac cell membrane.



**Fig. 9.** Detection of mRNAs coding for four isoforms of mAChRs (m<sub>1</sub>, m<sub>2</sub>, m<sub>3</sub>, and m<sub>4</sub>) in RNA samples extracted from canine atrium. Shown is a representative gel with ethidium bromide staining of PCR-amplified products. Lane 1, 100-bp cDNA size maker; lanes 2 through 9, expected PCR products for m<sub>1</sub> (338-bp band), m<sub>2</sub> (297-bp band), m<sub>3</sub> (216-bp band), and m<sub>4</sub> (257-bp band), respectively; RT+, reaction with reverse transcriptase; and RT-, omission of reverse transcriptase from the reaction for negative control.

First, our patch-clamp studies, using TMA and 4AP as pharmacological probes and various mAChR subtype-selective antagonists, provide for the first time strong functional evidence for cardiac M<sub>3</sub> and M<sub>4</sub> receptors. The concentrations of antagonists used in our experiments were chosen based on previous studies in the literature and the data from our binding assays, which should provide optimal selectivity to each subtype. For positive control, we also tested the effects of Meth on ACh-induced K<sup>+</sup> current. The fact that Meth (20 nM) indeed inhibited I<sub>KACH</sub> argued for the validity of concentrations used in our study. On the other hand, the drastic effects of M<sub>3</sub>-selective inhibitors could hardly be explained by cross-actions on other subtypes of mAChRs because we did not observe any changes in I<sub>KACH</sub> and I<sub>K4AP</sub> after the application of M<sub>3</sub> antagonists. Moreover, the significant shifts in concentration-response curves of I<sub>KTMA</sub> and I<sub>K4AP</sub> produced specifically by M<sub>3</sub> and M<sub>4</sub> antagonists, respectively, provide more conclusive evidence. The receptor protection-inactivation experiments further strengthen the evidence for the presence of M<sub>3</sub> receptors. The fact that PZ, an M<sub>1</sub>-selective antagonist, failed to alter I<sub>KTMA</sub> and I<sub>K4AP</sub> rules out the contribution of M<sub>1</sub> receptors to these two currents. The discrepancy between the study of Sharma et al. (1996) and ours likely represents the interspecies difference or tissue type heterogeneity, or both, considering that the rat ventricular cells were used in their study and canine atrial myocytes were used in the present study. The second line of evidence came from our receptor-binding assays of the displacement [<sup>3</sup>H]NMS binding by various mAChR antagonists. PZ, Meth, 4-DAMP, or Trop yielded pK<sub>i</sub> values pointing to the conclusion that there is coexistence of multiple subtypes of mAChRs (M<sub>2</sub>/M<sub>3</sub>/M<sub>4</sub>). The displacement of [<sup>3</sup>H]NMS binding by PZ yielded a high- and a low-affinity binding site with pK<sub>i</sub> values of 6.9 and 5.8, respectively. The high-affinity binding identifies M<sub>3</sub> and M<sub>4</sub> subtypes, and the value for the low-affinity binding is more suggestive of the presence of an M<sub>2</sub> subtype, which is inconsistent with its affinity to an M<sub>1</sub> receptor. Although the high-affinity binding of Meth (pK<sub>i</sub> = 7.7) suggests the existence of M<sub>2</sub>, M<sub>4</sub>, and M<sub>1</sub> subtypes, considering that PZ binding ruled out the possibility of M<sub>1</sub> in canine tissues, it is more likely that Meth actually binds to only M<sub>2</sub> and M<sub>4</sub> receptors (van Zwieten and Doods, 1995). The low-affinity binding of Meth (pK<sub>i</sub> = 6.6) identifies but does not distinguish M<sub>3</sub> and M<sub>5</sub> subpopulations. Similarly, 4-DAMP binding revealed two groups of mAChRs, with high-affinity binding (pK<sub>i</sub> = 9.1) in agreement with its affinity to M<sub>3</sub> and M<sub>1</sub> receptors and low-affinity binding (pK<sub>i</sub> = 7) for an M<sub>2</sub> subtype of mAChRs. However, in respect to PZ binding, the high-affinity pK<sub>i</sub> value for 4-DAMP binding would more likely correspond to an M<sub>3</sub> receptor. Competition binding of Trop gave a high-affinity binding site (pK<sub>i</sub> = 7.6) that fits well with the existence of an M<sub>4</sub> receptor subtype (Lazareno et al., 1990; Lazareno and Birdsall, 1993). The low-affinity binding of Trop (pK<sub>i</sub> = 6.1) does not seem to identify any subtypes of the known mAChRs. Taken together, the results from our binding experiments suggest the coexistence of multiple subtypes, M<sub>2</sub>, M<sub>3</sub>, M<sub>4</sub>, and probably M<sub>5</sub> as well, of mAChRs in the canine atrium. More interestingly, the K<sub>d</sub> value from binding assay is almost identical with the concentrations of 4-DAMP (2 nM) to block I<sub>KTMA</sub> and of Trop (200 nM) to block I<sub>K4AP</sub>. Finally, the fact that cDNA fragments representing m<sub>2</sub>, m<sub>3</sub>, and m<sub>4</sub> isoforms were isolated from

canine atrium and that expression of mRNAs coding for  $M_2$ ,  $M_3$ , and  $M_4$  subtypes was detected by RT-PCR adds an additional evidence for the presence of  $M_2/M_3/M_4$  mAChRs in the canine atrium.

Our conclusion relies largely on pharmacological dissection, and false interpretation could have been drawn due to imperfect specificity of mAChR antagonists currently available. Nevertheless, similar approach has been widely used for identifying and classifying mAChR subtypes. Moreover, our conclusion of multiple subtypes of mAChR was further supported by the results from molecular cloning and mRNA expression. No attempt was made to study  $m_5$  subtype in this work because there are no  $m_5$  subtype-selective antagonists available at this time and the potential physiological counterpart is still unknown.

mAChRs have been shown to coupled to several different ion channels. Interaction of  $M_2$  with  $I_{K_{ACh}}$  probably is the best known example. Other examples include  $M_2$  coupling to a nonselective cation current in smooth muscle cells (Wang et al., 1997) and  $m_4$  coupling to a cloned inward rectifier  $K^+$  channel, GIRK1 (Gadbut et al., 1996). We found here that both  $M_3$  and  $M_4$  are coupled to  $K^+$  channels that possess biophysical and pharmacological characteristics different from other  $K^+$  currents. Neither of the currents are sensitive to dofetilide, indapamide, and 293B, suggesting minimal contribution of the classic delayed rectifier  $K^+$  currents  $I_{K_r}$  and  $I_{K_s}$  (Wang et al., 1994). Both currents are completely suppressed by atropine and the mAChR subtype-selective antagonists, effects requiring activation of mAChRs. The two channels conduct both outward currents with characteristics of delayed rectifier currents on depolarization and inward currents similar to inward rectifier  $K^+$  currents in response to hyperpolarizing pulses. It is possible that these currents represent integral of two separate populations of channels: one conducting the delayed rectifier-like outward current and the other conducting the inward  $K^+$  current.  $I_{KTMA}$  and  $I_{K4AP}$  have many differences and similarities as well in terms of their voltage-dependence and kinetics. Whether they represent a single entity or two separate populations of channels is not clear at this time. Although further investigation is absolutely necessary for answering the questions raised above, the issue is beyond the scope of this study.

Our binding experiments showed that both TMA and 4AP were able to displace in a competitive manner the binding of [ $^3H$ ]NMS to mAChRs. The low-affinity  $K_d$  value of TMA binding (2.5 mM) is almost identical with the value (2.2 mM) for cloned  $m_3$  receptor reported by Wess et al. (1992). In addition, the  $K_d$  values of TMA and 4AP for receptor binding were well coincided with the concentrations of these compounds required for current induction (Table 2).

TMA has been reported to slow the sinus rate and to weaken the contraction of rat hearts (Zakharov et al., 1993). Our results revealed a possible mechanism underlying, at least in part, the TMA-produced negative inotropic and chronotropic effects. TMA activates  $I_{KTMA}$ , which should cause hyperpolarization of the membrane and shortening of the action potential duration. Hyperpolarization of membrane could lead to diminished automaticity, thereby slowing heart rate. On the other hand, shortening of action potential would indirectly decrease calcium entry into the cell, which can in turn result in reduction of contractile. 4AP has been reported to induce a delayed rectifier-like  $K^+$  current in isolated cat

atrial myocytes (Navarro-Polanco and Sánchez-Chapula, 1997), which was antagonized by atropine and probably mediated through a PTX-sensitive G protein, consistent with our findings in canine cells. In addition, although  $I_{K4AP}$  in dog has wave form and kinetics different from the 4AP-induced currents in cat, the two currents seem to share many similarities in terms of their pharmacological sensitivity and the voltage-dependent properties. It is quite likely that the feline 4AP-induced current is equivalent to  $I_{K4AP}$  in dog and also is mediated by an  $M_4$  receptor.

Our findings raised several questions regarding the  $M_3$  and  $M_4$  receptors in the heart. First, are there any endogenous activators specific for  $M_3/M_4$  subtypes? Our data in this study showed that ACh, in the presence of  $M_2$  antagonist Meth to inhibit  $M_2$  receptors, increased an outward current with properties distinct from  $I_{K_{ACh}}$  (Fig. 7). Similar effects were observed in 16 of 29 cells. Our preliminary data demonstrate that this outward current is significantly blocked by 2 nM 4-DAMP or 200 nM Trop, indicating the possible role of  $M_3$  or  $M_4$  receptors, or both. One major difficulty of dissecting this component from  $I_{K_{ACh}}$  is that Meth cannot completely block  $M_2$  without inhibiting  $M_3/M_4$  receptors. Moreover, ACh has been shown to inhibit the 4AP-induced  $K^+$  current (Navarro-Polanco and Sánchez-Chapula, 1997). We do not know at this time why ACh activates the time-dependent outward currents only when  $M_2$  receptors are largely blocked by the antagonists. Our data do not explain why only a certain percentage of cells possesses the outward current (in the presence of ACh and Meth), and we are unable to provide more reliable evidence for the activation of  $I_{KTMA}/I_{K4AP}$  by ACh. Many factors might be involved in this effect, including possible allosteric action of ACh binding to the receptors, heterologous desensitization between different subtypes of mAChRs, or possibly other unknown mechanisms. More careful and detailed investigation is absolutely needed to clarify this issue. We have also obtained data indicating that choline, a precursor and metabolite of ACh, is able to activate currents identical with  $I_{KTMA}$ , and the action of choline is also mediated by  $M_3$ -subtype receptors (unpublished observations). Second, what are the physiological roles of these novel subtypes of mAChRs and their coupled  $K^+$  channels in cardiac functions? We have performed preliminary experiments and acquired data showing that  $M_3$  activation produced significant slowing of the heart rate and shortening of the action potential duration. Obviously, further investigations are warranted to establish the roles of the cardiac  $M_3$  and  $M_4$  subtypes and their coupled  $K^+$  currents. Third, how are  $M_3$  and  $M_4$  receptors coupled to their  $K^+$  channels? Are they coupled directly via G proteins and other membrane-delimited components or indirectly through the cytoplasmic factors? Although it is of great importance to have better understanding of this aspect, it is out of the scope of the present investigation.

In summary, we have obtained strong functional and molecular evidence for the presence of multiple subtypes of mAChRs ( $M_2/M_3/M_4$ ) as well as the physiological functions of these receptor subtypes in the canine heart. The studies described have the potential to affect greatly our thinking about parasympathetic control of the heart. If the presence and importance of  $M_3$  and  $M_4$  receptors are confirmed, we will no longer be able to consider parasympathetic effects on the heart as due to a simple ACh/ $M_2$  interaction. These

findings potentially open up new opportunities for novel insight into the parasympathetic control of heart functions and the interactions between receptor and channel proteins.

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