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Molecular interactions between general anesthetics and the $5 H T_{\mbox{\tiny 2B}}$ receptor

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Abstract

Background: Serotonin modulates many processes through a family of seven serotonin receptors. However, no studies have screened for interactions between general anesthetics currently in clinical use and serotonergic G-protein-coupled receptors (GPCRs). Given that both intravenous and inhalational anesthetics have been shown to target other classes of GPCRs, we hypothesized that general anesthetics might interact directly with some serotonin receptors and thus modify their function.

Methods: Radioligand binding assays were performed to screen serotonin receptors for interactions with propofol and isoflurane as well as for affinity determinations. Docking calculations using the crystal structure of 5-HT_{2B} were performed to computationally confirm the binding assay results and locate anesthetic binding sites.

Results: The 5-HT_{2B} class of receptors interacted significantly with both propofol and isoflurane in the primary screen. The affinities for isoflurane and propofol were determined to be 7.78 and . 95 μ M, respectively, which were at or below the clinical concentrations for both anesthetics. The estimated free energy derived from docking calculations for propofol (–6.70 kcal/mol) and isoflurane (–5.10 kcal/mol) correlated with affinities from the binding assay. The anesthetics were predicted to dock at a pharmacologically relevant binding site of 5HT_{2B}.

Conclusions: The molecular interactions between propofol and isoflurane with the 5-HT_{2B} class of receptors were discovered and characterized. This finding implicates the serotonergic GPCRs as potential anesthetic targets.

Keywords

isoflurane; propofol; anesthetics; serotonin receptor

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Introduction

Serotonin (5-hydroxytryptamine; 5-HT) is an endogenous monoamine neurotransmitter involved in many central nervous system (CNS) processes, including the modulation of behavior, mood, aggression, anxiety, and nociception – an area of natural interest for anesthesiology (Bradley et al., 1986; Hamon et al., 1989; Kesim et al., 2005; Roth, Ciaranello, & Meltzer, 1992). Serotonin mediates its actions through a family of receptors divided into seven different classes and located both in the CNS and the periphery. Though six of these classes are composed of G-protein-coupled receptors (GPCRs), past research on the interactions between anesthetic agents and the serotonergic receptors has primarily focused on the 5-HT₃ class, a group comprising ligand-gated ion channel receptors. These receptors structurally and functionally resemble the heavily studied neuronal nicotinic acetylcholine receptors, GABA receptors, and glycine receptors (Joshi et al., 2006). This focus has resulted in numerous electrophysiological studies that have examined the modulatory effects of many general anesthetics vis-à-vis the 5-HT₃ class of receptors (Barann et al., 2000; Barann, Linden, Witten, & Urban, 2008; Jenkins, Franks, & Lieb, 1996; Machu & Harris, 1994; Suzuki, Koyama, Sugimoto, Uchida, & Mashimo, 2002; Zhang, Oz, Stewart, Peoples, & Weight, 1997).

Despite the focus on the 5-HT₃ receptors, anesthetic-related research outside of this class has been comparatively sparse and focused on the effects of anesthetics on the downstream physiological endpoints modulated by the serotonergic system. Previous studies have demonstrated that anesthetics at millimolar concentrations interact with two types of serotonin receptors (5-HT_{2A} and 5-HT₃) (Barann et al., 2000; Minami, Minami, & Harris, 1997). Despite these early findings and mounting evidence demonstrating that both intravenous and volatile anesthetics directly interact with GPCRs, no studies have systematically determined whether any of the other 5-HT receptors interact with general anesthetic agents at pharmacologically relevant concentrations (Hollmann, Strumper, Herroeder, & Durieux, 2005; Ishizawa, Pidikiti, Liebman, & Eckenhoff, 2002; Peterlin, Ishizawa, Araneda, Eckenhoff, & Firestein, 2005).

Given that GPCRs are anesthetic targets and that some of the physiological endpoints mediated by the serotonergic receptors overlap with side effects from anesthetic treatments, in the present study, we searched for possible interactions between the 5-HT receptors and two different anesthetics commonly utilized in surgical procedures – propofol (intravenous) and isoflurane (inhalational). We hypothesized that general anesthetics might interact directly with some serotonin receptors and thus modify their function through either agonist or antagonist modes of action. We screened available serotonin receptors with these two common clinical general anesthetics and the candidate receptor was further investigated through the use of high-resolution crystal structure data.

Materials and methods

Isoflurane and propofol were obtained from Halocarbon Laboratories (Liberty Corner, NJ). All other chemicals were of reagent grade or better and obtained from Sigma–Aldrich (St. Louis, MO).

Radioligand binding assays

A primary binding assay with 11 different 5-HT receptors was performed to determine the class of receptors for which the anesthetics displayed potential interaction. Evidence for interaction was based on the inhibition of the reference ligand-binding signal. Propofol, isoflurane, and the reference ligands for each receptor were diluted in standard binding buffer (50 mM Tris-HCl, 10 mM MgCl₂, .1 mM EDTA, and pH 7.4) to a final concentration of 10 μ M. The radioligands used were [³H] Way100635 at a final concentration of .4nM for 5-HT_{1A}; [³H] 5-CT at final concentrations in the range of 1.5–2.0 nM for 5-HT_{1B} and 5-HT_{1D}; [³H] 5-HT at a final concentration of 3.5 nM for 5-HT_{1E}; [³H] Ketanserin at a final concentration of 1.5 nM for 5-HT_{2A}; [³H] LSD at final concentrations in the range of 1–5 nM for 5-HT_{2B}, 5-HT_{5A}, 5-HT₆, and 5-HT_{7A}; [³H] Mesulergine at a final concentration of 3 nM for 5-HT_{2C}; and [³H] GR65630 at a final concentration of 1 nM for 5-HT₃. Fifty microliter aliquots of the radioligand were added to wells of a 96-well plate, which contained 25 µl aliquots of references or test ligands. Crude membrane fractions containing the receptors were suspended in standard binding buffer and were added to each well. Detailed information about membrane preparation can be obtained from the protocol online (http://pdsp.med.unc.edu/PDSP%20Protocols%20II%202013-03-28.pdf, pp. 18–24, last date accessed: 15 October 2013). The reactions were incubated at room temperature for 1.5 h to allow radioligand binding equilibration. Bound radioactivity was harvested by rapid filtration onto a .3% polyethyleneimine-treated 96-well filter mat using a 96-well filtermate harvester. The dried filters were treated with melted scintillant and a Microbeta scintillation counter was used to measure the radioactivity retained on the filter. Quadruplicate samples corresponding to every ligand concentration point were counted.

The secondary binding assay was performed only when a test compound had the signal inhibition greater than 50%. A secondary binding assay was performed to determine the binding affinity for each respective anesthetic. Propofol, isoflurane, and methysergide (reference ligand) were prepared in the standard binding buffer and serially diluted to the following concentrations (5× of final concentrations): 0 nM, .05 nM, .5 nM, 1.5 nM, 5 nM, 15 nM, 50 nM, 150 nM, 500 nM, 1.5 μ M, 5 μ M, and 50 μ M.

 $50 \ \mu\text{L}$ aliquots of [³H] LSD (5nM) were added to wells of a 96-well plate, which contained 25 μ l aliquots of the reference or test ligands. Fifty microliter of a crude membrane fraction of cells expressing the 5-HT_{2B} were added to each well. Reaction incubation, harvesting, and radioactivity measurement procedures from the primary assay were repeated. Triplicate samples corresponding to every ligand concentration point were counted.

Docking

Structures of isoflurane (CID: 3763), propofol (CID: 4943), dopamine (CID: 681), serotonin (CID: 5202), DOI (4-iodo-2,5-dimethoxyphenylisopropylamine, CID: 1229), LSD (lysergic acid diethylamide, CID: 5761), methysergide (CID: 9681), and ketanserin (CID: 3822) were obtained from the PubChem compound database. These structures were appropriately protonated and energy minimized using the PRODRG (Schüttelkopf & van Aalten, 2004) server. Then the ligand input files for docking were prepared using AutoDock Tools by removing non-polar hydrogen and setting up active torsions. The same tool was used to

prepare the receptor input file by adding polar hydrogen to the crystal structure of 5-HT_{2B} (PDB: 4IB4 (Wacker et al., 2013)).

AutoDock Vina (Trott & Olson, 2009) was used to perform docking of the above eight ligands into 5-HT_{2B}. In all cases, the side chains of the receptor were considered rigid. The search volume was selected to cover the extracellular oriented "half" of 5-HT_{2B} with dimensions 25 Å \times 25 Å \times 25 Å. The global search consisted of the default eight independent runs starting from distinct random positions of the ligand relative to the receptor. For each ligand, only the top-scored binding pose was considered. Due to the non-deterministic nature of the search algorithm, 100 dockings were repetitively performed with different initial random seeds. Then the most popular binding mode was selected, and the estimated binding affinities for structures with this mode were averaged for each ligand.

Data analysis

The data are presented as mean \pm SE from quadruplicate results. The results were analyzed using GraphPad Prism (version 5.02 Windows version, GraphPad Software Inco, La Jolla, CA).

Results

Primary binding assay

Among the 11 different receptors tested in the primary binding assay, only the 5-HT_{2B} subclass of serotonin receptors demonstrated significant signal inhibition in the presence of 10 μ M of propofol and isoflurane (Table 1). Propofol but not isoflurane treatment significantly inhibited the binding signal in 5-HT_{5A}. Further investigation on this receptor was not pursued because the secondary binding assay confirmed that the affinity would be higher than 10 μ M if there is any specific interaction.

Affinity determination

In the secondary assay, binding by both anesthetics occurred in a concentration-dependent manner comparable to the 5-HT_{2B} antagonist methysergide (Figure 1). The values of (K_i) for isoflurane (7.78 µM) and propofol (.95 µM) were substantially higher than that of methysergide (19 nM), indicating lower affinity.

Docking

The affinities estimated from the docking experiments correlated closely to the experimentally derived affinities (Figure 2). The calculated affinities for propofol (-6.7 kcal/mol) and isoflurane (-5.10kcal/mol) were comparable to those of the endogenous neurotransmitters dopamine (-5.90kcal/mol) and serotonin (-6.50 kcal/mol) as well as the agonist 2,5-dimethoxy-4-iodoamphetamine (DOI, -6.30 kcal/mol). The estimated affinities for the anesthetics were substantially weaker than those of the agonist lysergic acid diethylamide (LSD, -10.20 kcal/mol) and the antagonist ketanserin (-10.50 kcal/mol).

Based on the docking results, the putative binding pocket was mainly defined by residues from transmembrane 3 (TM3), TM5, TM6, and TM7. Asp^{3.32} hydrogen bonds with Tyr^{7.43}

(both residues are strictly conserved among all serotonin receptors, opioid receptors, adrenergic receptors, and histamine receptors). The disruption of this hydrogen bond by a charge–charge interaction between $Asp^{3.32}$ and the amine moiety of the agonists is believed to trigger the subsequent conformation change of the helices bundle from an inactive state to active state (Nichols & Nichols, 2008). The van der Waals volumes of ligands were compatible with the 5-HT_{2B} binding pocket (Figure 3(a) and (b)). Among them, the antagonist ketanserin is the largest ligand and occupies the entire putative binding pocket.

Asp^{3.32} is a highly conserved residue amongst the serotonergic receptors and it is believed to act as a counter-ion for the protonated amine of various ligands (Kristiansen et al., 2000; Wang, Gallaher, & Shin, 1993). The interaction between Asp^{3.32} and the amine group of methysergide, serotonin, dopamine, DOI, LSD, and ketanserin was also observed in the docking results. Among them, the distance between the amine hydrogen and the carboxylic oxygen of Asp^{3.32} was 2–3 Å for methysergide, serotonin, DOI, LSD, and dopamine.

Another residue, Ser^{3.36}, provides a second interaction site for the protonated amine of serotonin but not of LSD or bufotenin (Almaula, Ebersole, Zhang, Weinstein, & Sealfon, 1996). This interaction was also observed in our docking results. The distance between the amine hydrogen and the hydroxyl oxygen of Ser^{3.36} was 2.6 Å for both DOI and dopamine, respectively. However, this interaction distance is much larger for methysergide and LSD: 4.2–5.0 Å.

Propofol and isoflurane lack this amine moiety and therefore bind to 5-HT_{2B} in different ways. Phe^{6.52} and Trp^{6.48} are believed to be involved in anchoring the aromatic moiety of the ligands (Roth, Shoham, Choudhary, & Khan, 1997b). The binding poses in the docking showed that most ligands (except isoflurane) placed their aromatic rings close to these aromatic residues. Accordingly, propofol bound with 5HT_{2B} through van der Waals interactions between its phenyl ring and the aromatic residues $\text{Trp}^{6.48}$, Phe^{6.51}, and Phe^{6.52} in TM6 of 5HT_{2B} (Figure 3(b)).

Isoflurane does not possess an aromatic group, and the docking results showed that hydrogen bonding plays an important role in its interaction with $5HT_{2B}$ (Figure 3(c)). The main determinant of binding of isoflurane to $5HT_{2B}$ appeared to be hydrogen bonding between its ethanol oxygen and Thr^{3.37} (with a O–O distance of 3.1 Å). A fluorine atom in the trifluoromethyl group is poised to hydrogen bond with this same residue (with a F–O distance of 3.4 Å). A fluorine atom in the difluoro methyl moiety on the other side of the ligand was poised to form a hydrogen bond with Ser^{3.36} (with a F–O distance of 4.3 Å).

Discussion

A radioligand binding screen on the serotonin receptors identified interactions between general anesthetics with the $5HT_{2B}$ subclass of receptors. Secondary binding assays confirmed the interactions and determined that the affinities were below commonly used clinical concentrations. The results from the docking calculations on the crystal structure of $5HT_{2B}$ (PDB: 4IB4) are consistent with the experimental data and provide insights into how anesthetics interact with $5HT_{2B}$ receptors.

GPCRs are an important area of research for anesthesiology as the signaling cascades for many members of these membrane proteins modulate nociception both in the CNS and the periphery. Various agents commonly used in the perioperative period such as opiates, α_2 -adrenergic agonists, adenosine, and other derivative compounds have long been recognized as direct molecular targets for GPCRs (Hollmann et al., 2005). Moreover, in recent years, mounting evidence has suggested that inhalational anesthetics such as halothane and isoflurane either directly target or indirectly modify the functionality of these targets (Icaza et al., 2009; Ishizawa et al., 2002; Peterlin et al., 2005).

Studies exploring and characterizing potential interactions between general anesthetics and all the serotonergic GPCRs had previously been lacking, but the primary binding assay demonstrated that both propofol (intravenous) and isoflurane (inhalational) exhibited significant interactions with 5-HT_{2B} due to the large signal inhibition (Table 1). Propofol but not isoflurane exhibited potential interaction with 5-HT_{5A}, a GPCR whose physiological functions are relatively obscure though some data implicate it in the control of circadian rhythms (Thomas, 2006). However, a secondary binding study failed to demonstrate any specific interactions. Both isoflurane and propofol failed to demonstrate significant binding with the ligand-gated ion channel 5-HT₃ receptor in the primary assay. Though other studies have reported the potentiation of this receptor following isoflurane treatment, the lack of significant binding signal demonstrated here may be explained by the low concentration utilized in our study (10 μ M). The concentrations of isoflurane used in these previously studies were several orders of magnitude higher than that used in the current study (Barann et al., 2000; Jenkins et al., 1996; Suzuki et al., 2002). The receptor's lack of interaction with propofol correlated with previous electrophysiological studies which demonstrated that it had no effect on channel potentiation (Machu & Harris, 1994; Zhang et al., 1997).

Results from the secondary binding assay determined that the binding affinity (K_i) of isoflurane and propofol for 5-HT_{2B} are 7.78 and .95 µM, respectively. While the affinity constants for the anesthetics deviate from the classical values for specific interactions, these results remain relevant as both affinity constants represent values that are at or below the clinically effective concentrations for both compounds. Because propofol binds to soluble human blood proteins, thereby leaving 1–3% of the compound free for potential interactions with receptors, its clinically effective concentration is generally considered to be ~1 µM (Franks & Lieb, 1994; Servin, Farinotti, Haberer, & Desmonts, 1993). The free aqueous concentration of isoflurane needed for anesthesia is often reported to be 100–300 µM (Franks, 2008; Franks & Lieb, 1993, 1998).

The docking position and pose of methysergide overlaps with that of ergotamine, which is present in the crystal structure (Figure 4(a)) (PDB: 4IB4). The protonated amines of methysergide and ergotamine align very well and they both interact with the carboxylic oxygen of $Asp^{3.32}$ (H–O distances 2.1 and 1.8 Å, respectively). Almaula et al. found that mesulergine (selective for 5-HT_{2C}) and ergoline (selective for 5-HT_{2A}) reversed their relative affinity with mutations A5.46S for 5-HT_{2C} and S5.46A for 5HT_{2A} (Almaula, Ebersole, Ballesteros, Weinstein, & Sealfon, 1996). Thus, they proposed that a hydrogenbonding interaction existed between Ser^{5.46} of 5-HT_{2A} and the N1 hydrogen of N1-unsubstituted ergolines. At the same position, Ala^{5.46} of 5-HT_{2C} facilitates the interaction

with N1 methyl group of N1-substituted ergolines. Our docking results were consistent with this, showing a distance of 3.0 Å (C–C distance) between the N1 methyl of methysergide and Ala^{5.46} of 5-HT_{2B}. LSD, which has the same binding pose as methysergide but contains N1 hydrogen instead, had a slightly lower binding affinity to 5-HT_{2B} (.1 kcal/mol difference). The docking results showed that the hydroxyl group of serotonin is in proximity to and poised for hydrogen bond with Ser^{5.43} with a H–O distance of 2.1 Å, which might explain previous observations (Johnson, Wainscott, Lucaites, Baez, & Nelson, 1997) for 5-HT_{2A} where serotonin showed over a 10-fold decrease of affinity with the mutation S5.43A. This interaction with Ser^{5.43} was also observed for one hydroxyl group of dopamine in our docking results. Docking calculations for isoflurane, propofol, and methysergide yielded binding affinities that correlated with the experimental results (Figure 2).

The docking position of propofol also overlaps with that of ergotamine in the crystal structure (Figure 4(b)) (PDB: 4IB4). Propofol binding was partly caused by interactions between the molecule's phenyl ring and the aromatic residues $Trp^{6.48}$, $Phe^{6.51}$, and $Phe^{6.52}$. Interestingly, this aromatic cluster in helix VI is conserved between the three $5HT_2$ receptor subtypes. Previous work with $5HT_{2C}$ models proposed that $Trp^{6.48}$ and $Phe^{6.52}$ were involved in the binding of serotonin through the formation of hydrophobic interactions with the phenyl rings of these ligands (Kristiansen & Dahl, 1996). Mutagenesis studies with $5HT_{2A}$ further confirmed the importance of interactions with these residues for agonist, but not antagonist, binding (Choudhary, Craigo, & Roth, 1993; Roth, Shoham, Choudhary, & Khan, 1997a). Furthermore, $Phe^{6.52}$ has been reported to play a role in agonist activation of the $5HT_{2A}$ receptor (Braden, Parrish, Naylor, & Nichols, 2006).

Studies on the contributions of Phe^{6.51} interaction, however, have provided conflicting results. Early work indicated that the residue did not participate in agonist binding but rather contributed to antagonist stabilization (Choudhary et al., 1993; Roth et al., 1997a). However, recent evidence has suggested that it is involved in agonist binding (Braden et al., 2006). Based on these previous findings and our results, the mode of action of propofol on the $5HT_{2B}$ receptor cannot be extrapolated and further testing will be needed to determine this. However, our results nonetheless suggest that propofol docks at a binding site with pharmacological activity as the interacting aromatic residues on helix VI mediate G-protein activation and second-messenger production via direct coupling to the third intracellular loop (Suel, Lockless, Wall, & Ranganathan, 2003; Wess, Brann, & Bonner, 1989).

Our results demonstrated that hydrogen bonding interactions with Ser^{3.36} and Thr^{3.37} were important for isoflurane binding with the receptor. Much like propofol, isoflurane's mode of action on the receptor cannot be extrapolated, but its interactions occur at or near a pharmacologically relevant site as it overlaps with the one for the natural endogenous ligand serotonin.

The $5HT_{2B}$ receptors have been implicated in mediating several downstream physiological endpoints. In the CNS, the receptors are responsible for modulating mood, behavior, migraines, and nociception. In the periphery, the receptor has been found in arterial endothelium and the characterization of its cardiovascular effects has been a major area of research (Kaumann & Levy, 2006; Tanaka, Ludwig, Kersten, Pagel, & Warltier, 2004). In

particular, the subclinical value of the affinity constant coupled with our computational studies of ligand–receptor interactions could further elucidate the mechanisms underlying the side effects of propofol treatment.

Decreases in systemic arterial pressure commonly occur following anesthetic administration (Boer, Bovill, Ros, & van Ommen, 1991; Claeys, Gepts, & Camu, 1988; Rouby et al., 1991; Turner et al., 1998). Multiple mechanisms have been proposed to explain this phenomenon. Some studies have pointed to an inhibition of sympathetic vasoconstrictive nerve activity (Hoka, Yamaura, Takenaka, & Takahashi, 1998; Robinson, Ebert, O'Brien, Colinco, & Muzi, 1997). Others have implicated myocardial depression (Pagel & Warltier, 1993) while considerable evidence has suggested a nitric oxide-mediated mechanism and therefore a direct anesthetic action on smooth muscle relaxation (Gragasin & Davidge, 2009; Muzi, Berens, Kampine, & Ebert, 1992; Petros, Bogle, & Pearson, 1993; Rouby et al., 1991). Nitric oxide (NO) produced and released from the vascular endothelium plays a key role in vascular dilatation (Palmer, Ashton, & Moncada, 1988). It mediates systemic, coronary, and pulmonary vascular tone through a guanylyl cyclase-dependent mechanism in vascular smooth muscle cells (Levin, 1995). Evidence has implicated the activation of $5HT_{2B}$ in the production of NO in the endothelial cells of the human coronary artery and consequent arterial relaxation (Ishida, Kawashima, Hirata, & Yokoyama, 1998). Further evidence has also demonstrated that 5HT_{2B} is the receptor responsible for modulating endothelialdependent vasodilation in porcine pulmonary arteries (Glusa & Pertz, 2000; Glusa & Roos, 1996). Given the fact that 5HT_{2B} exists throughout the systemic vascular system (Choi & Maroteaux, 1996), the interaction between 5HT2B and propofol may explain the decreased systemic arterial pressure seen in patients during propofol and isoflurane administration. However, more in vitro and in vivo studies are needed to confirm such a hypothesis.

In conclusion, the molecular interactions between propofol and isoflurane with the 5-HT_{2B} class of receptors were discovered and characterized. This finding implicates the serotonergic GPCR as potential anesthetic targets.

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Figure 1.

Isoflurane and propofol binding affinity. Propofol exhibited higher affinity to 5-HT_{2B} compared to isoflurane; both values were lower than that of methysergide (5-HT_{2B} antagonist).



Figure 2.

Comparison between docking and experimental binding constants of isoflurane (I), propofol (P), and methysergide (M) to 5-HT_{2B}.





Figure 3.

(a) The van der Waals shapes of the ligands are compatible with the binding pocket.Propofol (magenta) docking within the pocket overlaps well with that of methysergide(cyan). (b) Stick representation of key residues in the binding pocket of 5-HT2B involved in propofol binding. (c) Isoflurane.

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Figure 4.

(a) Stick representation of overlapping between methysergide (cyan and docking result) and ergotamine (green, crystal structure). (b) Propofol (magenta) docking within the pocket overlaps well with that of ergotamine (green).

Table 1.

Signal inhibition of serotonin receptors by propofol and isoflurane.

Receptor	Isoflurane signal inhibition (%)	Propofol signal inhibition (%)
5-HT1A	26.4 ± 4.9	17.4 ± 6.6
5-HT1B	5.9 ± 2.3	12.7 ± 4.0
5-HT1D	3 ± 3.5	-6.1 ± 3.9
5-HT1E	-4.9 ± 2.0	-3.8 ± 3.0
5-HT2A	4.4 ± 7.4	7.2 ± 7.6
5-HT2B	61.7 ± 2.8	86.9 ± 1.1
5-HT2C	11.1 ± 5.8	48.8 ± 1.7
5-HT3	14.1 ± 6.2	28.4 ± 3.9
5-HT5A	54.2 ± 1.4	43.2 ± 2.0
5-HT6	30.8 ± 1.7	23 ± 2.2
5-HT7	$42.6\pm.9$	35.0 ± 2.6

Note: Data are presented as mean \pm SE from four replicates. Both anesthetics had a final concentration of 10 μ M when incubated with the receptors. Secondary binding assay for affinity determination was performed only when signal inhibition was greater than 50% for a given anesthetic.