

Pharmacologic analysis of non-synonymous coding 5-HT2A SNPs reveals alterations psychedelic drug potencies and efficacies

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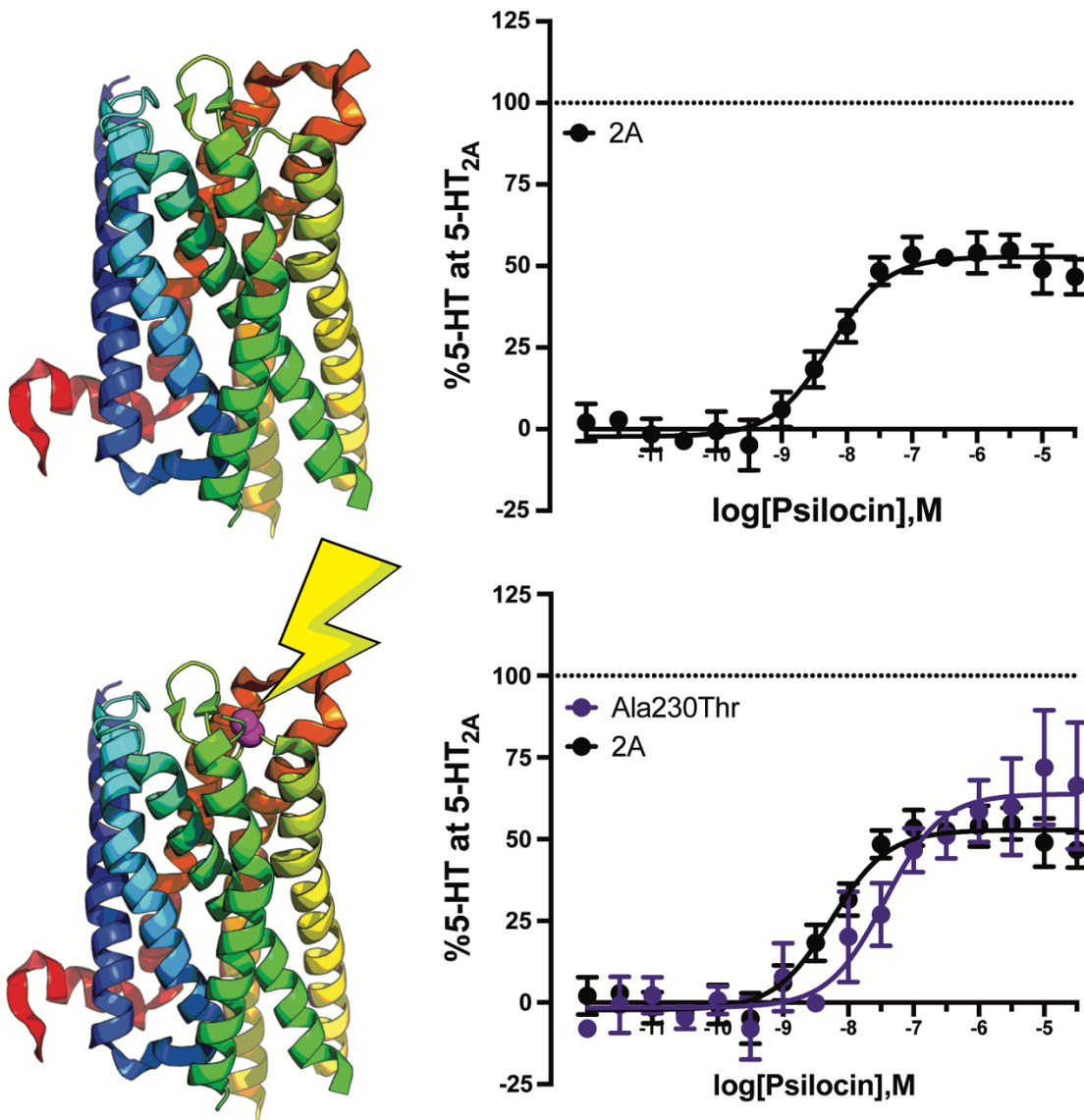
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Abstract

Serotonin (5-Hydroxytryptamine; 5-HT) 2A receptor (5-HT_{2A}R) signaling is essential for the actions of classical psychedelic drugs. In this study, we examined whether random sequence variations in the gene (single nucleotide polymorphisms, SNPs) encoding the 5-HT_{2A}R affect the signaling of four commonly used psychedelic drugs. We examined the *in vitro* pharmacology of seven non-synonymous SNPs, which give rise to S12N, T25N, D48N, I197V, A230T, A447V, and H452Y variant 5-HT_{2A} serotonin receptors. We found that these non-synonymous SNPs exert statistically significant, although modest, effects on the efficacy and potency of four therapeutically relevant hallucinogens. Significantly, the *in vitro* pharmacological effects of the SNPs drug actions at 5-HT_{2A}R are drug specific.

Keywords: 5-HT_{2A}, Serotonin Receptors, Hallucinogen, Psychedelic, SNP, Polymorphism

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1. Introduction

Psychedelic drugs have been defined as compounds that induce lysergic acid diethylamide (LSD)-like effects on cognition, emotion, and perception via activation of 5-HT_{2A} serotonin (5-hydroxytryptamine; 5-HT) receptors^{1, 2}. Recently, there has been renewed interest in psychedelic compounds as potential therapeutics for many neuropsychiatric disorders³. For example, psilocybin, the prodrug to the active compound psilocin found in *Psilocybe cubensis* mushrooms, has recently been granted breakthrough drug status by the Food and Drug Administration due to its potential as a treatment for treatment-resistant depression and anxiety^{4, 5, 6}. These effects of psilocybin in these Phase II clinical trials are both rapid and apparently enduring^{4, 5, 7}. Similarly, LSD has demonstrated efficacy in treating cluster headaches⁸ and alleviating anxiety in terminally ill patients⁹. Finally, anecdotal reports and recent meta-analyses have suggested the potential therapeutic utility for mescaline and 5-methoxy-*N,N*-dimethyltryptamine (5-MeO-DMT) in treating anxiety, depression, and other affective disorders^{10-12, 13, 14}.

Psychedelics are generally classified by chemical structure (e.g. tryptamines; ergolines and phenylisopropylamines) and each psychedelic drug has a robust pharmacology, with activities at many serotonin and other biogenic amine receptors¹⁵⁻¹⁹. 5-HT_{2A} receptor agonism has been shown to be crucial in mediating the psychoactive effects of psychedelics in both animal²⁰⁻²² and human studies²³⁻²⁶. 5-HT_{2A} receptors are members of the G protein-coupled receptors (GPCR) superfamily and canonically signal through the G_{αq} family of heterotrimeric G proteins activating phospholipase C_β and many other down-stream effector systems^{27, 28, 2}. 5-HT_{2A} receptors also recruit β-arrestin (βArr) proteins *in vitro*²⁹ and the 5-HT_{2A}R is apparently complexed with βArr1 and βArr2 in cortical neurons *in vivo*³⁰. While many GPCR agonists activate both G protein- and βArr-mediated signaling pathways, certain ligands are known to preferentially activate one signaling pathway via a ligand-dependent phenomenon known as functional selectivity or biased signaling³¹. It has been proposed that ligands with biased profiles may be useful for activating desired pathways (i.e. therapeutically

efficacious) rather than undesired pathways (i.e. unwanted side effect producing) downstream of a given GPCR³². Indeed, LSD is shown to preferentially signal through β Arr2 at the 5-HT_{2A} receptor^{18, 33} and numerous LSD-elicited responses are either significantly attenuated or completely absent in β Arr2-KO mice³⁴.

Early clinical studies on psilocin have shown a wide variability in drug response³⁵ with a significant portion of respondents showing no statistically significant treatment effect. The reasons for

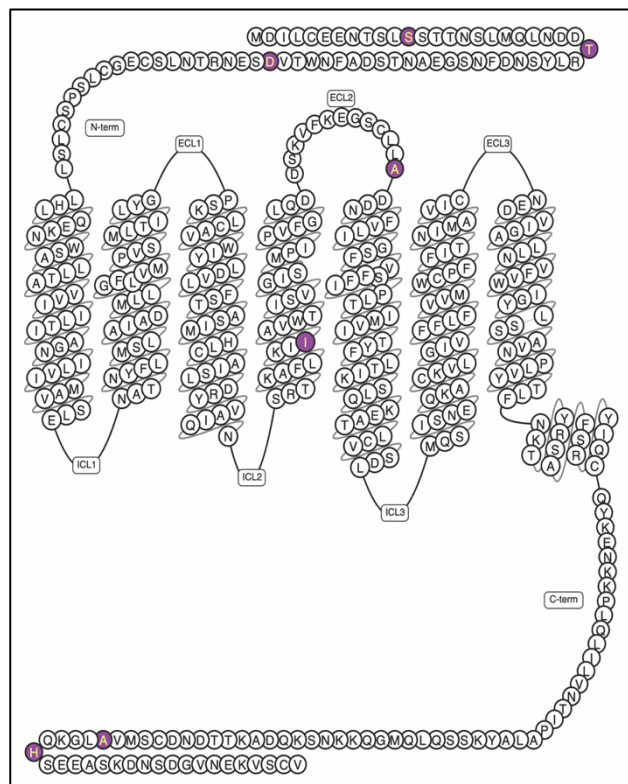


Figure 1: Snake plot representation of the 5-HT_{2A} receptor showing the location of the seven SNPs: S12N, T25N, D48N, I197V, A230T, A447V, H452Y.

such inter-individual variability in hallucinogenic drug response are unknown. At the molecular level, random sequence variations in genes (single nucleotide polymorphisms, SNPs) could explain inter-individual differences in drug response and such activities are highly relevant as hallucinogens become more prevalent in clinical practice. At least seven non-synonymous SNPs are located within the coding region of the human 5-HT_{2A} receptor gene (HTR2A, 13q14-21) (Fig. 1). These seven single-nucleotide polymorphisms (S12N, T25N, D48N, I197V, A230T, A447V, and H452Y) have potential actions on both receptor structure and function³⁶. As shown in Figure

1, the S12N, T25N, and D48N SNPs map to the predicted N-terminal tail of the receptor; the I197V SNP resides in the fourth transmembrane helix; the A230T SNP resides in the putative second extracellular loop (ECL2); the two remaining non-synonymous SNPs (A447V and H452Y) are located within the putative C-terminal tail. These SNPs represent seven of the most common variants observed in human populations with allele frequencies varying from 0.003% (T25N) to 7.9% (H452Y) (<https://gpcrdb.org/>)³⁷.

Here, we comprehensively examined the potency and signaling bias for four therapeutically promising psychedelics at several 5-HT_{2A} non-synonymous SNPs. We discovered that the effects of individual SNPs were drug selective and that no single SNP had uniform effects on all psychedelics tested. These results imply that future clinical studies into the therapeutic utility of psychedelics might consider particular drug-SNPs combinations – paying particular attention to those SNPs that cause significant differences in the *in vitro* pharmacologies of test agents.

2. Materials and Methods

2.1 Cell culture

HEK293T cells were obtained from ATCC (Manassas, VA). Cells were maintained, passaged, and transfected in Dulbecco's Modified Eagle's Medium containing 10% fetal bovine serum (FBS), 100 Units/mL penicillin, and 100µg/mL streptomycin (Gibco-ThermoFisher, Waltham, MA) in a humidified atmosphere at 37°C and 5% CO₂. After transfection, cells were plated in DMEM containing 1% dialyzed FBS, 100 Units/mL penicillin, and 100µg/mL streptomycin for BRET assays.

2.2 Mutagenesis

The human 5-HT_{2A} cDNA cloned into the *NotI* site pIRES-neo (Clontech) was obtained through the resources of the National Institute of Mental Health Psychoactive Drug Screening Program (<http://kidb.pdsp.unc.edu>). The following mutagenesis primers were obtained from Eton Bioscience:

S12N forward: 5' TCACTGAACAGCACAATACTCTCTCATG 3'

S12N reverse: 5' TGTGCTGTTTCAGTGAGGTATTCTCCTC 3'

T25N forward: 5' GATGATAACCGACTTTACTCCAACGACTTCAAC 3'

T25N reverse: 5' AAGTCGGTTATCATCATTGAGCTGCATGAG 3'

D48N forward: 5'- ACAGTAAACAGTGAAAACCCFACCAATCTGTCC

D48N reverse: 5'- TTCACTGTTTACTGTCCAATTGAAGGCGTC

I197V forward: 5' AAAATCGTCGCGGTCTGGACCATTTC 3'

I197V reverse: 5' GACCGCGACGATTTTCAGAAAGGCTTT 3'

A230T forward: 5' CTTCTGACTGATGACAATTCGTACTTATAGGAAGC 3'

A230T reverse: 5' GTCATCAGTCAGAAGGCAACTGCCCTC 3'

A447V forward: 5' ATGGTGGTCCTGGGCAAACAGCACAGTGAAGAAGCC 3'

A447V reverse: 5' GCCCAGGACCACCATACTGCAGTCGTTGTC 3'

H452Y forward: 5' AAACAGTACAGTGAAGAAGCCTCCAAAGAC 3'

H452Y reverse: 5' TTCACTGTACTGTTTGCCCAGGGCCAC 3'

SNPs were introduced into the cloned human 5-HT_{2A} receptor template using the QuikChange site directed, PCR-based mutagenesis kit exactly as described by the manufacturer (Stratagene, La Jolla, CA, USA). Mutagenized clones were isolated and subjected to Sanger DNA sequencing to verify the entire coding sequence for the presence of the desired mutation and the absence of any PCR-induced sequence errors.

2.3 BRET2 assays

Cells were plated either in six-well dishes at a density of 700,000–800,000 cells/well, or 10-cm dishes at 7–8 million cells/dish. Cells were transfected 2–4 hours later, using a 1:1:1:1 DNA ratio of receptor:Gα-RLuc8:Gβ:Gγ-GFP2 (100 ng/construct for six-well dishes, 1000 ng/construct for 10-cm dishes). Transit 2020 (Mirus Biosciences, Madison, WI) was used to complex the DNA at a ratio of 3 μL Transit/μg DNA, in OptiMEM (Gibco-ThermoFisher, Waltham, MA) at a concentration of 10 ng DNA/μL OptiMEM. The next day, cells were harvested from the plate and plated in poly-D-lysine-coated white, clear bottom 96-well assay plates (Greiner Bio-One, Monroe, NC) at a density of 30,000–50,000 cells/well.

One day after plating in 96-well assay plates, white backings (Perkin Elmer, Waltham, MA) were applied to the plate bottoms, and growth medium was carefully aspirated and replaced immediately with 50 μL of assay buffer (1x HBSS + 20 mM HEPES, pH 7.4) containing freshly prepared 50 μM coelenterazine 400a (Nanolight Technologies, Pinetop, AZ). After a five-minute equilibration period, cells were treated with 50 μL of drug (2X) for an additional 5 minutes. Plates were then read in a Pherastar FSX microplate reader (BMG Labtech Inc., Cary, NC) with 395 nm (RLuc8-coelenterazine 400a) and 510 nm (GFP2) emission filters, at 1 second/well integration times. BRET2 ratios were computed as the ratio of the GFP2 emission to RLuc8 emission. Results are from at least three

independent experiments, each performed in duplicate. Data were normalized to 5-HT stimulation and analyzed using nonlinear regression “log(agonist) vs. response” in Prism 9 (Graphpad Software Inc., San Diego, CA).

2.4 Bioluminescence Resonance Energy Transfer (BRET) Arrestin Assay

To measure 5-HT_{2A}R-mediated β -arrestin1 recruitment, HEK293T cells were co-transfected in a 1:1:5 ratio with human 5-HT_{2A}R containing C-terminal *Renilla* luciferase (*RLuc8*), GRK2, and Venus-tagged N-terminal β -arrestin1. To measure 5-HT_{2A}R-mediated β -arrestin2 recruitment, HEK293T cells were co-transfected in a 1:1:5 ratio with human 5-HT_{2A}R containing C-terminal *Renilla* luciferase (*RLuc8*), GRK2, and Venus-tagged N-terminal β -arrestin2. For both experiments, after at least 24 hours, transfected cells were plated in poly-lysine coated 96-well white clear bottom cell culture plates in plating medium (DMEM + 1% dialyzed FBS) at a density of 40,000–50,000 cells in 200 μ L per well and incubated overnight. The next day, medium was decanted, and cells were washed with 50 μ L of drug buffer (1X HBSS, 20 mM HEPES, 0.1% BSA, 0.01% ascorbic acid, pH 7.4), then 50 μ L of drug buffer with *RLuc* substrate coelenterazine h (Promega, 5 μ M final concentration) was added per well. After 5 minutes, 50 μ L of drug (2X) was added per well and incubated for 5 minutes. Plates were read for both luminescence at 485 nm and fluorescent eYFP emission at 530 nm for 1 second per well using a Pherastar FSX microplate reader (BMG Labtech Inc., Cary, NC). The ratio of eYFP/*RLuc* was calculated per well and the net BRET ratio was calculated by subtracting the eYFP/*RLuc* per well from the eYFP/*RLuc* ratio in wells without Venus- β -Arrestin present. Results are from at least three independent experiments, each performed in duplicate. The net BRET ratio was plotted as a function of drug concentration using Prism 9 (Graphpad Software Inc., San Diego, CA). Data were normalized to 5-HT stimulation and analyzed using nonlinear regression “log(agonist) vs. response” in Prism 9.

2.5 Receptor Expression Measurements

HEK293T cells were plated in 10cm dishes and transfected with the wild-type 5-HT_{2A} R, S12N, T25N, D48N, I197V, A230T, A447V, or H452Y variant of 5-HT_{2A} receptor. After 36h of transfection, cell

medium was replaced with medium containing 1% dialyzed FBS. After 48 h, medium was aspirated and cells were washed twice with cold phosphate-buffered saline (PBS). Cells were scraped in 10 mL ice cold PBS and centrifuged at 3000 rpm at 4°C for 15 minutes and the supernatant was discarded. The cell pellet was resuspended in ice cold lysis buffer (50 mM Tris HCl, pH 7.4) and incubated on ice for 20 min. After incubation, cells were triturated gently for hypotonic lysis. The suspension was centrifuged at 21000g for 20 minutes at 4°C to obtain a crude membrane pellet. The supernatant was discarded, and the membrane pellets were stored at -80°C for further use.

Saturation binding assays were performed to determine equilibrium dissociation constant (K_d) and B_{max} . Membrane pellets were dissolved in lysis buffer and the amount of protein was estimated using Bradford assays. Binding assays were performed in standard binding buffer (50 mM Tris, 10 mM $MgCl_2$, 0.1 mM EDTA, 0.1% BSA, 0.01% ascorbic acid, pH 7.4) using [3H] ketanserin (PerkinElmer, Waltham, MA; specific activity 22.8 Ci/mmol) as the radioligand. Saturation binding assays were carried out in 96-well plates in a final volume of 125 μ L per well. In brief, 25 μ L of radioligand was added to a 96-well plate followed by addition of 25 μ L binding buffer (for total binding) or 25 μ L reference compound (Clozapine) at a final concentration of 10 μ M to measure nonspecific binding. After, 75 μ L of fresh membrane protein (15 μ g/well) was added and plates were incubated in the dark at room temperature for 90 minutes. The reaction was stopped by vacuum filtration onto cold 0.3% polyethyleneimine (PEI)-soaked 96-well filter mats using a 96-well Packard Filtermate harvester, followed by three washes with cold wash buffer (50 mM Tris, pH 7.4). Filters were dried and scintillation cocktail (Meltilex, PerkinElmer) was melted onto the dried filters and allowed to cool to room temperature. Then, filters were placed into cassettes and radioactivity was measured in a Microbeta counter (Molecular Devices). Data were analyzed using “One Site-Total and nonspecific binding” equation using Prism 9. Specific binding was calculated after subtracting the non-specific binding.

2.6 Data Analysis

All concentration-response curves were fit to a three-parameter logistic equation in Prism (Graphpad Software, San Diego, CA). BRET2 concentration-response curves were analyzed as either raw net BRET2 (fit Emax-fit Baseline) or by normalizing to a reference agonist for each experiment. Efficacy (Emax) calculations were performed according to Kenakin and colleagues³⁸: stimulus-response amplitudes (net BRET2) were normalized to the maximal responding agonist (maximal system response). EC50 and Emax values were estimated from the simultaneous fitting of all biological replicates. Transduction coefficients were calculated as $\text{Log}(\text{Emax} / \text{EC50})$ as described in Kenakin and colleagues³⁸ and propagation of error was conducted at all steps. EC50, Emax, and transduction coefficient values were analyzed first by ANOVAs (F-test of curve fit, one-way ANOVA, or two-way ANOVA as described in the text). *Post hoc* pair-wise comparisons used Tukey-adjusted *p* values to control for multiple comparisons. Significance threshold was set at $\alpha = 0.05$.

3.0 Results

3.1 G_{α_q} protein mediated signaling activity potencies for therapeutically promising psychedelics differ between wild-type and polymorphic 5-HT_{2A} receptors:

To determine whether naturally occurring, non-synonymous SNPs in the 5-HT_{2A} receptor are likely to affect psychedelic actions *in vivo*, we measured the *in vitro* potencies and efficacies of the wild-type, S12N, T25N, D48N, I197V, A230T, A447V, and H452Y variant 5-HT_{2A} receptors for four therapeutically promising psychedelics. As shown in Figure 2 and Table 1, certain polymorphic 5-HT_{2A} receptors exhibited statistically significant, albeit modest, changes in response to specific psychedelics. The Ala230Thr and His452Tyr receptors displayed a sevenfold decrease in potency for

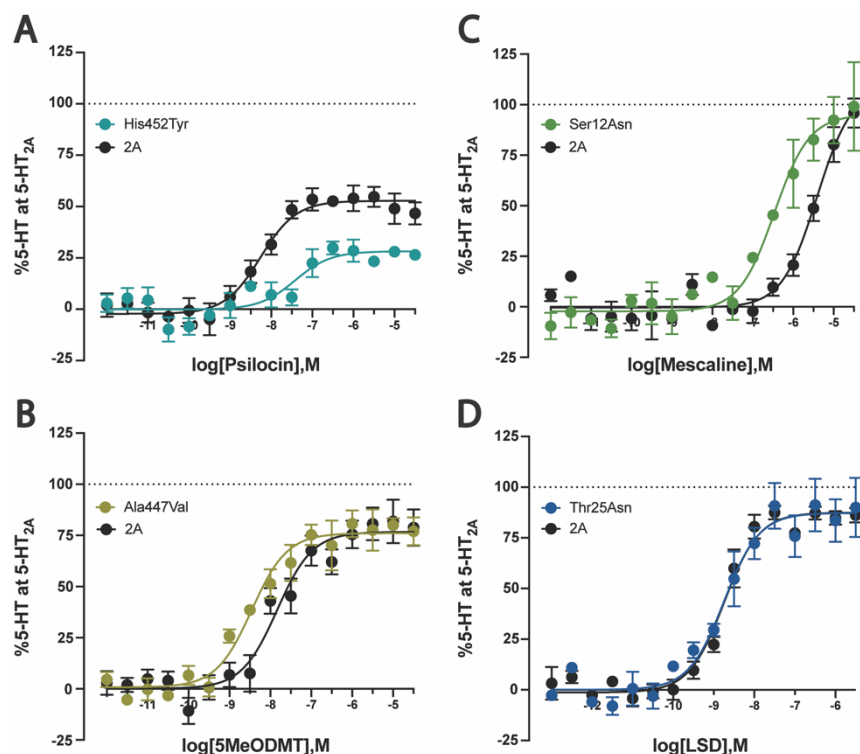


Figure 2: Representative Gq dis concentration-response curves for psychedelic drugs (a) Psilocin, (b) LSD, (c) Mescaline, and (d) 5-MeO-DMT at wild-type and polymorphic 5-HT_{2A} receptors using TRUPATH BRET assays. See Table 1 for the complete data matrix.

Drug	pEC50 ± s.e. (EC50, nM)							
	WT	Ser12Asn	Thr25Asn	Asp48Asn	Ile197Val	Ala230Thr	Ala447Val	His452Tyr
Psilocin	-8.261±0.123 (5.48)	-7.98±0.132 (10.47)	-8.126±0.088 (7.48)	-8.034±0.119 (9.25)	-8.223±0.113 (5.99)	-7.418±0.299* (38.2)	8.409±0.087 (3.90)	-7.419±0.247* (38.2)
5-MeODMT	-7.975±0.135 (10.6)	7.608±0.20 8 (24.7)	-8.246±0.155 (5.67)	-7.820±0.117 (15.2)	-7.795±0.132 (16.0)	-8.399±0.301 (4.00)	-8.481±0.191* (3.30)	-8.442±0.226 (3.61)
Mescaline	-5.413±0.108 (3860)	-6.468±0.127*** (340)	-6.060±0.120* (871)	-6.149±0.153*** (710)	-5.678±0.145 (2100)	-5.775±0.211 (1680)	-6.025±0.170** (943)	-5.577±0.159 (2650)
LSD	-8.732±0.074 (1.85)	8.600±0.12 2 (2.51)	-8.773±0.129 (1.69)	-8.862±0.111 (1.37)	-8.615±0.102 (2.43)	-8.716±0.225 (1.92)	-8.769±0.076 (1.70)	-8.441±0.192 (3.63)

Table 1: Potency (pEC50) values for several potentially therapeutic psychedelics at wild-type (WT) and polymorphic 5-HT_{2A} receptors as measured using TRUPATH Gq BRET assays. * $P < 0.05$, ** $P < 0.005$, *** $P < 0.0005$ by F-test.

psilocin compared to wild type (38nM vs 5nM, $P<0.05$). The Ala447Val 5-HT_{2A} receptor displayed a threefold increase in potency for 5-MeO-DMT compared to the wild type (3.3nM vs 10.6nM, $P<0.05$). Four polymorphic 5-HT_{2A} receptors exhibited statistically significant changes in mescaline potency. Of these, the largest effect was observed for the Ser12Asn 5-HT_{2A} receptor, which displayed a tenfold increase in potency for mescaline compared to the wild type (340.4 nM vs 3863 nM, $P<0.0005$). None of the SNP-induced alterations in LSD potency were significantly different from the wild-type value ($P<0.05$).

3.2 β Arrestin mediated signaling activity affinities for therapeutically promising psychedelics differ between wild-type and polymorphic 5-HT_{2A} receptors:

Having addressed whether several therapeutically promising psychedelics exhibit differential *in vitro* G α_q protein signaling pharmacologies at wild-type and polymorphic 5-HT_{2A} receptors, we next investigated effects on β Arrestin recruitment. As shown in Figure 3 and Table 2, the β Arr1 potencies of all psychedelics tested were not significantly altered by any of the SNP 5-HT_{2A} receptors

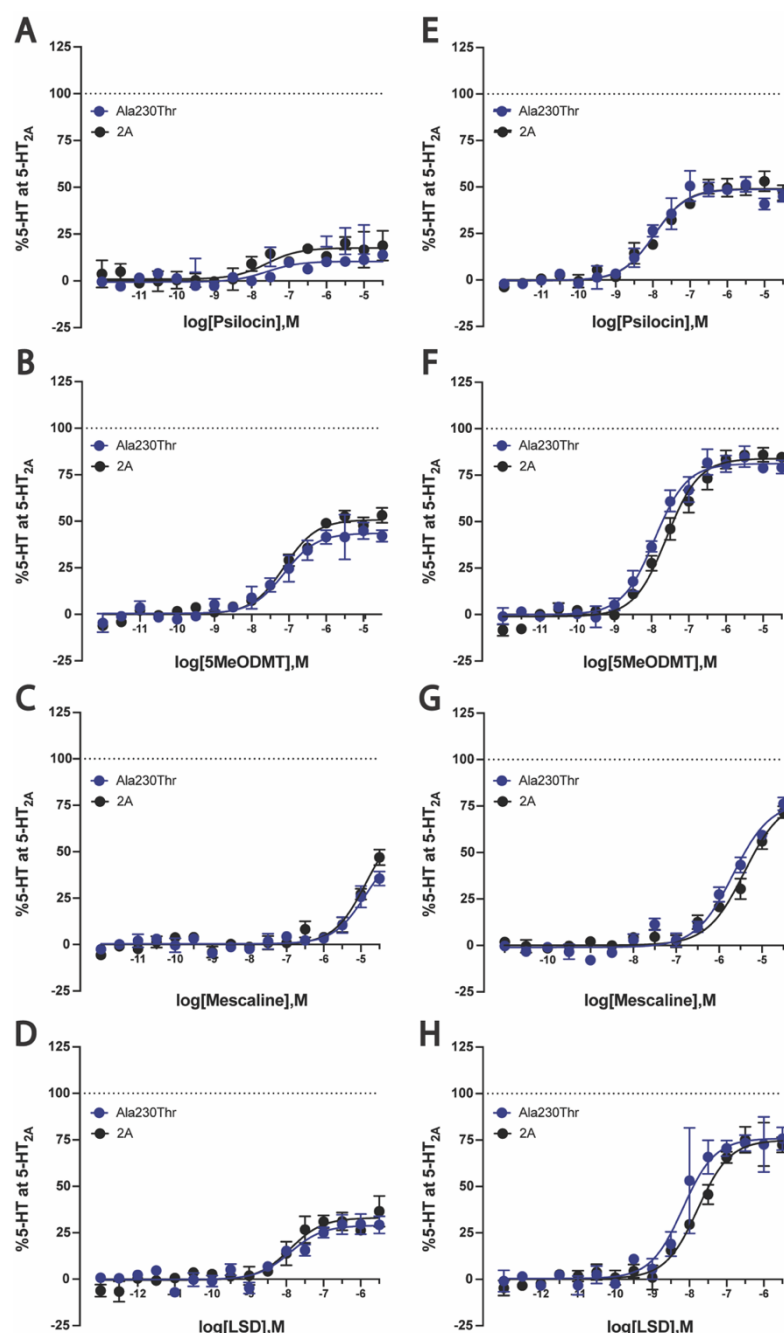


Figure 3: Concentration-response curves for psychedelic drugs (a) Psilocin, (b) LSD, (c) Mescaline, and (d) 5-MeODMT at wild-type and polymorphic Ala230Thr 5-HT_{2A} receptors using BRET assays of β arrestin-1 and -2 recruitment. See Tables 2 and 3 for the complete data matrix.

tested. In contrast, the β Arr2 potencies of 5-MeO-DMT, mescaline, and LSD tested at the Ala230Thr 5-HT_{2A} receptor were significantly altered. Specifically, the 5-MeO-DMT potency was increased twofold compared to wild type, mescaline potency was increased twofold compared to wild type, and LSD potency was increased threefold compared to wild type. As shown in Figure 3 and Table 3, the Ala230Thr 5-HT_{2A} receptor was the only SNP found to cause significant changes in β Arrestin2 potencies of 5MeODMT, LSD, and mescaline.

Drug	pEC50 \pm s.e. (EC50, nM)							
	WT	Ser12Asn	Thr25Asn	Asp48Asn	Ile197Val	Ala230Thr	Ala447Val	His452Tyr
Psilocin	-7.853 \pm 0.352 (14.04)	-6.907 \pm 0.351 (123.9)	-7.631 \pm 0.375 (23.38)	-7.681 \pm 0.302 (20.84)	-8.090 \pm 0.302 (8.13)	-7.101 \pm 0.390 (79.21)	-7.385 \pm 0.256 (41.20)	-7.726 \pm 0.328 (18.81)
5-MeODMT	-7.088 \pm 0.073 (81.68)	-7.169 \pm 0.190 (67.75)	-7.073 \pm 0.177 (84.57)	-7.154 \pm 0.123 (70.22)	-7.078 \pm 0.170 (83.48)	-7.222 \pm 0.144 (59.95)	-7.363 \pm 0.325 (43.32)	-7.105 \pm 0.142 (78.44)
Mescaline	-4.786 \pm 0.162 (16365)	-5.421 \pm 0.350 (3797)	-4.226 \pm 0.576 (59436)	-5.296 \pm 0.211 (5057)	-5.062 \pm 0.304 (8669)	-5.023 \pm 0.167 (9494)	-4.926 \pm 0.334 (11869)	-5.408 \pm 0.243 (3905)
LSD	-7.942 \pm 0.175 (11.42)	-7.742 \pm 0.214 (18.10)	-7.643 \pm 0.186 (22.74)	-7.813 \pm 0.172 (15.37)	-7.780 \pm 0.182 (16.61)	-7.793 \pm 0.164 (16.12)	-8.050 \pm 0.255 (8.92)	-7.583 \pm 0.243 (26.12)

Table 2: Potency (pEC50) values for several potentially therapeutic psychedelics at wild-type (WT) and polymorphic 5-HT_{2A} receptors as measured using β Arr1 recruitment BRET assays.

Drug	pEC50 \pm s.e. (EC50, nM)							
	WT	Ser12Asn	Thr25Asn	Asp48Asn	Ile197Val	Ala230Thr	Ala447Val	His452Tyr
Psilocin	-7.812 \pm 0.094 (15.42)	-8.067 \pm 0.155 (8.57)	-7.802 \pm 0.120 (15.77)	-7.611 \pm 0.129 (24.50)	-7.878 \pm 0.094 (13.24)	-8.087 \pm 0.116 (8.18)	-7.810 \pm 0.121 (15.48)	-7.927 \pm 0.097 (11.84)
5-MeODMT	-7.595 \pm 0.064 (25.40)	-7.629 \pm 0.094 (23.51)	-7.624 \pm 0.074 (23.75)	-7.647 \pm 0.086 (22.56)	-7.707 \pm 0.088 (19.62)	-7.916\pm0.071** (12.13)	-7.482 \pm 0.127 (32.98)	-7.523 \pm 0.77 (30.00)
Mescaline	-5.410 \pm 0.089 (3887)	-5.266 \pm 0.118 (5421)	-5.383 \pm 0.105 (4143)	-5.343 \pm 0.096 (4537)	-5.363 \pm 0.112 (4333)	-5.675\pm0.076* (2114)	-5.258 \pm 0.154 (5527)	-5.324 \pm 0.076 (4748)
LSD	-7.781 \pm 0.086 (16.57)	-7.887 \pm 0.090 (12.97)	-7.924 \pm 0.063 (11.92)	-7.868 \pm 0.88 (13.56)	-7.720 \pm 0.062 (19.05)	-8.179\pm0.120* (6.62)	-7.755 \pm 0.150 (17.59)	-7.641 \pm 0.091 (22.84)

Table 3: Potency (pEC50) values for several potentially therapeutic psychedelics at wild-type (WT) and polymorphic 5-HT_{2A} receptors as measured using β Arr2 recruitment BRET assays. * P <0.05, ** P <0.005 by F-test.

3.4 Bias calculations for wild-type and polymorphic 5-HT_{2A} receptors

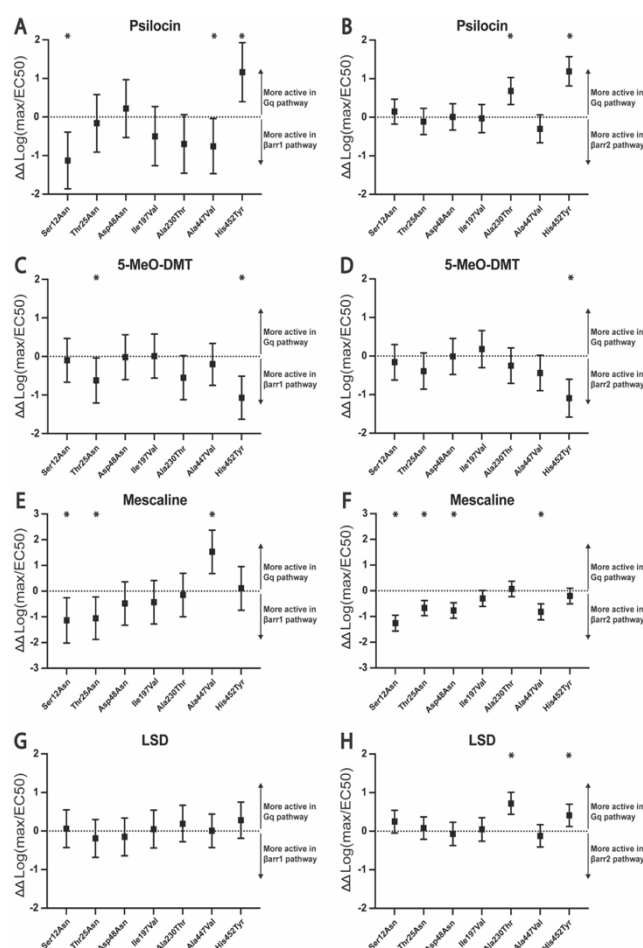


Figure 4: Bias plots showing $\Delta\Delta\log(\text{Emax}/\text{EC}_{50})$ values as a measure of ligand bias between signaling pathways. The error bars represent 95% confidence interval; when interval includes zero, the ligands are not biased with respect to each other. (A, B) Psilocin, (C, D) LSD, (E, F) 5-MeO-DMT, (G, H) Mescaline. (A, C, E, G) Gq versus $\beta\text{arr}1$, (B, D, F, H) Gq versus $\beta\text{arr}2$.

Having addressed whether several SNPs exhibit different *in vitro* signaling for several potentially therapeutic psychedelics, we next quantified changes in bias using the approach of Kenakin and colleagues³⁸. To compare between the wild-type and mutant receptors, $\Delta\Delta\log(\text{Emax}/\text{EC}_{50})$ values were calculated. As shown in Figure 4, overall biases [$\Delta\Delta\log(\text{Emax}/\text{EC}_{50})$ values] for the investigated pathways were significantly altered by multiple SNPs. Psilocin signaling was significantly altered by the Ser12Asn, Ala447Val, and His452Tyr SNPs. The His452Tyr 5-HT_{2A} receptor showed the greatest change in signaling, favoring G α_q signaling over $\beta\text{arr}1$ and $\beta\text{arr}2$. The His452Tyr 5-HT_{2A} receptor also demonstrated the largest change in 5-MeO-DMT signaling, increasing G α_q signaling relative to $\beta\text{arr}1$ and $\beta\text{arr}2$. Mescaline signaling was significantly

altered by the Ser12Asn, Thr25Asn, Asp48Asn, and Ala447Val 5-HT_{2A} receptors. Only the Ala230Thr and His452Tyr 5-HT_{2A} receptors demonstrated altered LSD signaling profiles, favoring G α_q over $\beta\text{arr}2$ in both cases.

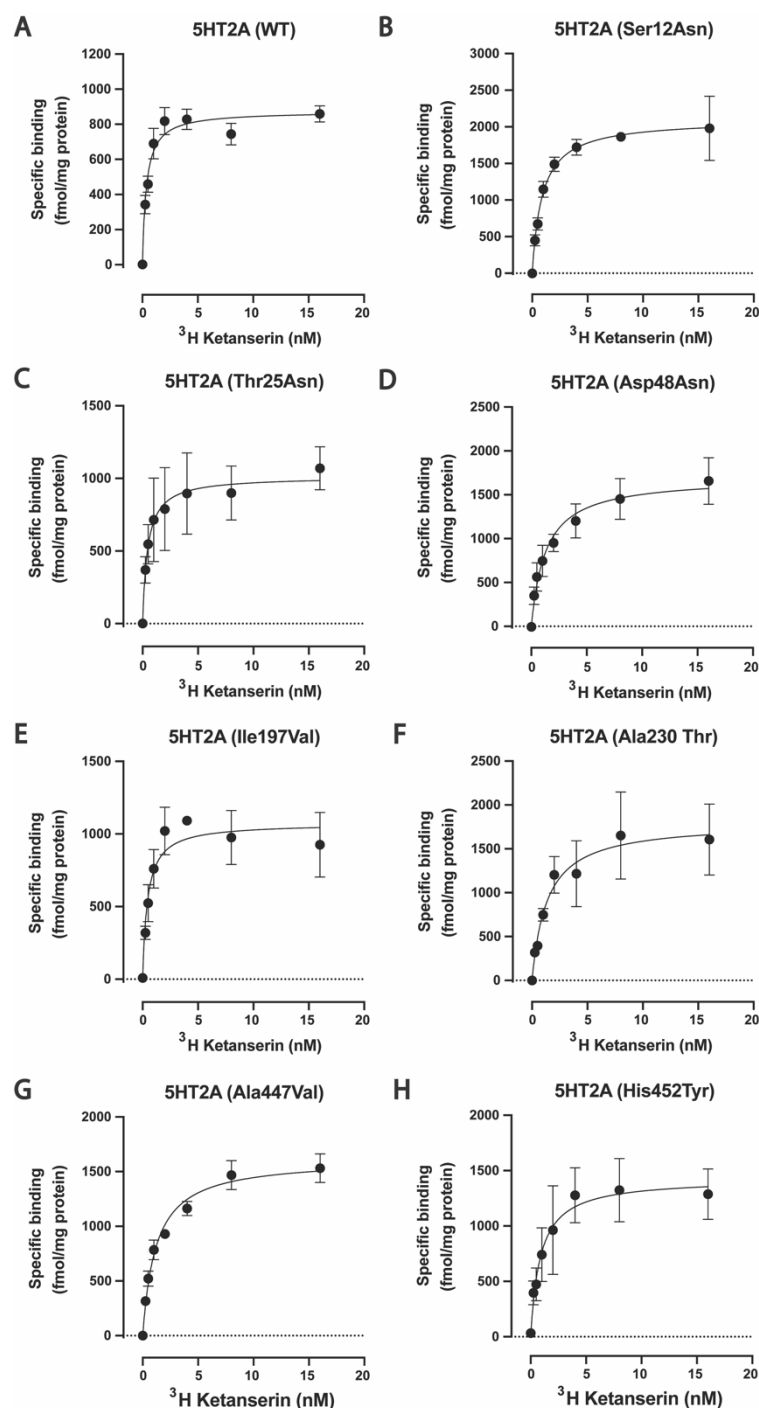


Figure 5: Saturation radioligand binding analysis of WT and polymorphic 5HT_{2A} receptors as a measure of receptor expression. (A) WT 5HT_{2A}, (B) Ser12Asn, (C) Thr25Asn, (D) Asp48Asn, (E) Ile197Val, (F) Ala230Thr, (G) Ala447Val, (H) His452Tyr.

3.5 Effects of 5-HT_{2A} receptor SNPs on receptor expression

To determine whether the 5-HT_{2A} receptor SNPs affected receptor expression, we measured the maximum binding (B_{max}) for each by radioligand saturation binding assay as shown in Figure 5 and Table 4. None of the SNPs altered receptor expression more than threefold compared to the wild type. The largest alteration of receptor expression was seen in the Ser12Asn 5-HT_{2A} receptor which was, on average, reduced less than threefold relative to the wild-type receptor. As we previously demonstrated our BRET-based assays are relatively insensitive to differences in receptor expression³⁹. We note that even though the Ser12Asn and Ala230Thr polymorphisms were both expressed at levels higher than WT no uniform differences in agonist potencies or efficacies were observed.

Thus, the differences observed between the

in vitro pharmacology of the seven variant 5-HT_{2A} receptors tested probably do not arise due to differential expression.

	WT	Ser12Asn	Thr25Asn	Asp48Asn	Ile197Val	Ala230Thr	Ala447Val	His452Tyr
Bmax (fmol/mg)	874.4±38.4	2100±113	1013±120	1698±148	1073±91.1	1802±215	1618±72.4	1429±185

Table 4: Bmax values at wild type (WT) and polymorphic 5-HT_{2A} receptors as measured using radioligand competition binding

4.0 Discussion

The major finding of this study is that certain 5-HT_{2A} receptor non-synonymous SNPs can alter the *in vitro* pharmacology of the tested psychedelics. As we have demonstrated in the current study, even polymorphisms some distance from the orthosteric binding site of the receptor can have substantial effects on the *in vitro* pharmacology of a given drug. Given that several psychedelic drugs are in clinical trials, this information could aid in the design and ultimate interpretation of clinical studies.

Our prior study³⁶ examined some of these polymorphisms on the agonist and antagonist actions of several drugs, although functional selectivity was not. Additionally, of the compounds tested in the current study, only 5-MeO-DMT was examined. Based on the current results, our data imply that individuals carrying the Ala230Thr variant might show altered responses to all of the psychedelics studied given that G α_q and/or β arr2 signaling was altered. Additionally, we would predict that individuals carrying the His452Tyr variants might show altered responses to psilocin and psilocybin, as large decreases in psilocin G α_q potency were found to significantly alter the signaling bias. For mescaline, the most dramatic change was seen at the Ser12Asn polymorphism with a tenfold increase in potency. Of note, our study only examined binding potency and efficacy at a single five-minute timepoint. β arr2 signaling of LSD, for instance, has been shown to increase in potency and efficacy over time¹⁸. Therefore, future studies could examine the effects of 5-HT_{2A} receptor SNPs on signaling kinetics.

In summary, our findings indicate that 5-HT_{2A} receptor SNPs can alter the *in vitro* pharmacology of some therapeutically promising psychedelics. Prior studies have already demonstrated that the His452Tyr as well as Thr25Asn and Ile197Val 5-HT_{2A} receptor polymorphisms can exhibit pronounced differences from the wild-type receptor in the pharmacologies of atypical antipsychotics³⁶. Our results confirm that these receptors can exhibit pronounced differences from the wild-type receptor in the pharmacologies of psychedelics. Additionally, we find that the

polymorphisms Ser12Asn, Ala230Thr, and Ala447Val 5-HT_{2A} receptor polymorphisms also exhibit pronounced differences. These results may have relevance for the design and interpretation of future clinical trials.

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