



Experimental evaluation of the generalized vibrational theory of G protein-coupled receptor activation

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Recently, an alternative theory concerning the method by which olfactory proteins are activated has garnered attention. This theory proposes that the activation of olfactory G protein-coupled receptors occurs by an inelastic electron tunneling mechanism that is mediated through the presence of an agonist with an appropriate vibrational state to accept the inelastic portion of the tunneling electron's energy. In a recent series of papers, some suggestive theoretical evidence has been offered that this theory may be applied to nonolfactory G protein-coupled receptors (GPCRs), including those associated with the central nervous system (CNS). [Chee HK, June OS (2013) *Genomics Inform* 11(4):282–288; Chee HK, et al. (2015) *FEBS Lett* 589(4):548–552; Oh SJ (2012) *Genomics Inform* 10(2):128–132]. Herein, we test the viability of this idea, both by receptor affinity and receptor activation measured by calcium flux. This test was performed using a pair of well-characterized agonists for members of the 5-HT₂ class of serotonin receptors, 2,5-dimethoxy-4-iodoamphetamine (DOI) and *N,N*-dimethyllysergamide (DAM-57), and their respective deuterated isotopologues. No evidence was found that selective deuteration affected either the binding affinity or the activation by the selected ligands for the examined members of the 5-HT₂ receptor class.

protein | mechanism of action | pharmacology | quantum biology | electron tunneling

Olfaction—and other chemo-sensitive sensory processes—is an important information-gathering technique for organisms of many clades and kingdoms. Specifically, human olfaction is known to occur by activation of olfactory receptors (ORs)—a subclass of G protein-coupled receptors (GPCRs)—located within the nasal epithelium and mediating responses within the olfactory bulb, where it is encoded and conveys information to the amygdala, the orbitofrontal cortex, and the hippocampus (1, 2). The discovery and the cloning of ORs led to the joint 2004 Nobel Prize in Physiology and Medicine to Richard Axel and Linda Buck. (3) GPCRs are 7-helical transmembrane proteins, facilitating communication from extracellular ligand signals to the cellular interior through activation of (interior) G proteins, while maintaining the integrity of the membrane (4). GPCRs are activated by an appropriate agonist moving into the protein's orthosteric binding site, resulting in a conformational change within the helical bundle. This structural change leads to altered conformations of the intracellular loops that couple to appropriate signaling molecules within the cell, e.g., G proteins. A recent series of papers has experimentally determined activated/inactivated states through isotopic-tagged receptors with NMR spectroscopy (5–7). An additional work used molecular dynamics to provide structural insights into how the agonist may assist the interchange between conformations through several proposed peptide sidechain pathways by examining the structures of the activated and inactivated μ -opioid receptor (8). Additionally, photon-induced conformational changes in light-sensitive proteins have also been observed (9, 10).

Recently, an iteration of the vibrational theory of olfaction (VTO)—suggested and advocated by Luca Turin (11, 12)—

has arisen and has gained both supporters (13–16) and detractors (17, 18); the novelty of this incarnation of the VTO is ascribable to its nonthermal- and nonphoton-based mechanism. Turin's theory is a contemporary reincarnation of the more classical theory proposed by Dyson (19), Wright (20), and Wright and Serenius (21), where the activation of the olfactory receptor is performed—or sensitive to—the molecular vibrations of the olfactant. Dyson suggested that the molecular vibrations of the agonist were exactly responsible for the activation of the protein. These vibrations were entirely thermally excited, as no mechanism for photoexcitation is available within the body. The modern incarnation of this theory suggests that the OR behaves as an electron-tunneling (ET) junction. Hypothetically, this electron transfer instigates the conformation change of the olfactory GPCR, leading to the intracellular signal cascade. This theory both is contentious and lacks direct at-receptor evidence. Several of the proposed tests for this theory, including odor mixing, have been addressed to disprove this theory through controlled sensory tests (13, 15). Additionally, more modern examinations of this theory with both humans and insect subjects have provided mixed and controversial results. (14, 16–18, 22–24).

There have been several tests of the previous iterations of the VTO, including odor blending (25, 26) and isotope exchange (20, 27–30); each of these methods—odor blending (11, 18) and isotope exchange (11, 31)—has been used to evaluate the modern VTO. Initial justifications of the VTO correlated characteristics of the vibrational spectra with olfactory perception (12), continued within ORs (32), and expanded into non-OR GPCRs (33–36). Isotope exchange—which provides the ability to alter the vibrations of a molecule while maintaining its chemistry—may manifest itself as an alteration of either the intensity or the quality of the scent. Isotope exchange tests of the VTO focused on ¹H→²H exchange, although ¹³C exchange has been suggested

Significance

Herein, we test the present iteration of the vibrational theory of protein activation by comparing predictions obtained from Turin's vibrational theory for the activation of olfactory receptors measuring affinity and activation at a nonolfactory receptor family of G protein-coupled receptors. This was done at the CNS serotonin receptor family h5-HT₂ and with both the 2,5-dimethoxy-4-iodoamphetamine and *N,N*-dimethyllysergamide agonists. Invalidation was performed through a comparative analysis of agonist behavior between isotopologues.

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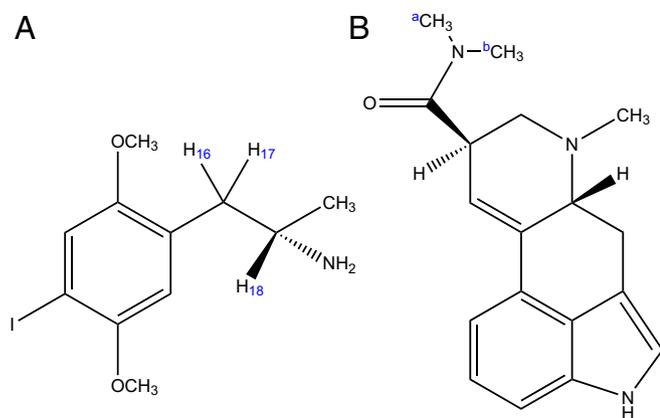


Fig. 1. (A) Structure of the (*R*)-DOI molecule. In blue text, the atomic indexes for some specific sidechain hydrogen atoms initially considered for deuteration are shown. Additional sites considered for deuteration are the hydrogens of the two methoxy groups. (B) Structure of the DAM-57 molecule. In blue text, each methyl amide carbon where deuteration exchange was undertaken is denoted as either carbon "a" or "b."

(37). Unfortunately, measuring either quality or intensity changes by sensory studies has presented itself as inconclusive and difficult to quantify.

Recently, several recent studies—both *in vitro* and *in vivo*—have been conducted at a more physiological level. Block et al. (17) studied the activation of several ORs (importantly, OR5AN1 and MOR244-3) by both muscone and cyclopentadecanone; neither receptor displayed a significant differential response to isotopologues of the musk odorants, suggesting failure of the VTO at mammalian ORs. However, criticism of this work has arisen on grounds of lacking the *in situ* environment (odorant binding proteins, cofactors, etc.) and possibly being too specific with respect to the repertoire of examined ORs (23, 24, 38, 39). Supporting previous behavioral studies (14, 16), Drimyl et al. (23) and Paoli et al. (24) studied direct electrophysical responses induced by odorant detection at the antennae and glomerular lobe of several *Drosophila* species and *Apis mellifera*, respectively. These findings show unique spatial-temporal responses with respect to families of isotopologues. Two things should be further noted with respect to these *in situ* insect studies: (i) Despite attempts at compensation, these results may not be entirely independent of perireceptor effects (including transport, enzymes, and extracellular vestibule). (ii) Insect ORs are ionotropic receptors (IRs) and not GPCRs; each class of receptors (GPCRs vs. IRs) shows specific evolutionary benefits (broad responsiveness vs. speed of detection and processing) (40, 41).

Herein, we exploit the fact that ORs are a subclass of the broader GPCR family with highly conserved sequences and structural motifs. The nonexceptionalism of ORs within the broader class of GPCRs was previously discussed within the context of the VTO (18) and is highlighted by Barwich (42), who asserts ORs are a model system for the GPCRs within neurobiology. Furthermore, ORs maintain up to 40% genetic similarity with rhodopsin (43). Additionally, ORs appear within areas of mammalian corpus that have no olfactory capacity (44–55). We therefore hypothesize that due to functional and morphological similarities, if ORs are activated through an ET mechanism, other GPCRs share the same fundamental mechanism. Examination of another (better characterized) GPCR subclass may provide insight into the possibility of the proposed ET mechanism. The serotonin 5-HT₂ receptor class, notably a primary target for hallucinogenic compounds, was selected as the main test for the contemporary VTO. (*R*)-2,5-dimethoxy-4-iodoamphetamine (*R*-DOI) is a well-characterized agonist for the serotonin 5-HT_{2A}

receptor used as a standard ligand for studying the pharmacology of 5-HT_{2A} receptors and is widely used as a radioligand to measure expression and affinity of ligands at 5-HT_{2A} receptors, particularly in brain tissues. (*R*)-DOI is reported to have psychedelic effects somewhat similar to those of lysergic acid diethylamide (LSD) (56). A quintessential hallucinogenic ligand for the 5-HT_{2A} receptor is LSD; a highly related (non-Schedule 1) molecule is generated by substituting the diethyl amide with a dimethyl amide, creating *N,N*-dimethyllysergamide (DAM-57). DAM-57 is an assumed serotonergic psychedelic with limited hallucinogenic capacity and very mild autonomic stimulation in humans at dosages of ~100 μg (57).

The organization of this paper is as follows: *Theoretical Predictions* gives a brief explanation of the theoretical predictions for (*R*)-DOI at members of the 5-HT₂ serotonin receptor class according to Turin's theory, while making direct reference to a previous work concerning similar predictions for DAM-57; *Experimental Procedures* briefly introduces methods of experimental analysis for determining affinity; *Results and Discussion* provides a discussion of the experimental results and a comparison with theoretical predictions; and finally, in *Conclusion* we provide concluding remarks.

Theoretical Predictions

Turin—within the contemporary VTO—hypothesized that the active site of the GPCR [specifically an OR, although later works considered generalizing this hypothesis (18, 36, 58)] acts as an ET junction (11). According to the theory, an electron emerges from a donation site—likely a metal atom acting as a cofactor (11, 31), redox chemistry (59), or peptide sidechain (11, 60) capable of oxidation—and traverses the active site to an acceptor site, which is likely a specific motif or residue sidechain. As the electron traverses the active site, it may undertake several paths: (i) elastic tunneling, where no energy is lost or gained by the electron; (ii) inelastic tunneling (IET), where the electron may donate or accept a quantum of energy during transfer; and (iii) subsequently higher ordered inelastic processes (61–63). The hypothesized presence of a possible metal cofactor site—acting to assist either in binding or in a later activation step—at ORs, GPCRs, and non-GPCR chemokine receptors is supported by altered behavioral response (64–66), physiological response (66–68), theory (69, 70), and *in vitro* observations (70–79).

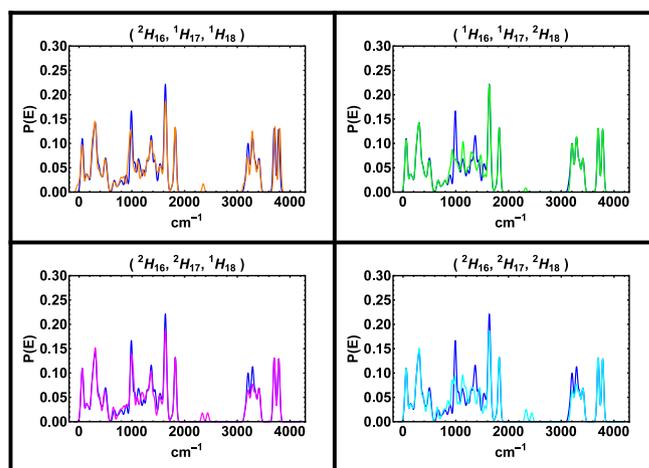


Fig. 2. Plots of the tunneling probability as a function of energy, $P(E)$, for various deuterated analogues of the DOI molecule. (Top Left) In orange, ²H₁₆ deuterated DOI; (Top Right) in green ²H₁₈ deuterated DOI; (Bottom Left) in magenta, ²H₁₆ and ²H₁₇ deuterated DOI; and (Bottom Right) in cyan, ²H₁₆, ²H₁₇, and ²H₁₈ deuterated DOI. In all plots, blue is the all-protium DOI tunneling spectrum.

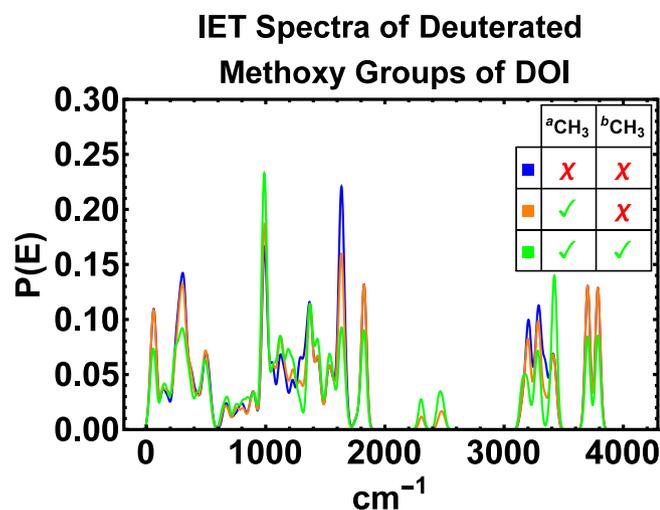


Fig. 3. Plots of the tunneling probability as a function of energy, $P(E)$, for various deuterated analogues of the (*R*)-DOI molecule with respect to the deuteration of one or both aromatic methoxy groups. Note that the double deuteration depletes the possible active peaks at 1,600–1,800 cm^{-1} by roughly 50%, indicating that this experiment could be carried out with a detectable effect.

Within the hypothesized protein-based ET junction, the donor and acceptor energies are offset by a quantum of energy, ΔE . If an electron attempts to undergo an elastic tunneling process, it has no available acceptor site. However, if the electron is capable of losing a specific quantum of energy (to an internal mode of a bound ligand) such that $E_{\text{acceptor}} - E_{\text{donor}} = \Delta E$, the transfer can be undertaken, activating the protein. Thus, a molecule must both fit into the active site with the correct orientation and have a vibrational mode capable of assisting in an IET process to activate the protein. Working within Turin's hypothesis, several theoretical expansions have been undertaken to account for specific considerations of the system, including charge transfer rates (60, 80), receptor effects (80–82), and chiral effects (83).

Block et al. (17) gave several criticisms of current modeling approaches to the VTO; these criticisms include inappropriate reorganization energies, not considering dynamic fluctuations on the system, excluding possible protein electron couplings during ET, and an unreliable electron delivery mechanism. Recently, Reese et al. (81) addressed several of these concerns, showing that the binding effects of the receptor have a nonzero—all but negligible—effect on the ligand, that dynamic fluctuations have a very small effect on the transfer, and that the reorganization energy (λ) for an OR can conform to $\lambda \ll 1$ kcal/mol. Concerns of electron density leaking into the environment were previously addressed through coupling a vibrational bath with the electron transfer (80), and such studies have shown that environmental-induced dissipation could enhance the vibrational signaling (82). Evaluating the reliability of the electron delivery mechanism cannot be addressed until the complete structure of the OR is known and models account for all cofactors including the possible effects of perireceptor molecular species, as there is evidence that NADPH and other oxidative processes are important in GPCR activation (59). Furthermore, future theoretical investigations should include the fundamental electron–dipole interaction—placing emphasis on the orientation of each vibrating dipole—described by Lambe and Jaklevic (61), Kirtley et al. (62), and Phillips and Adkins (63), among others. This interaction potential permits Raman modes to assist transfer and was recently used by Bittner et al. (84).

We—in the context of Turin's VTO—previously conducted a theoretical study of several ligands belonging to the family of

serotonin receptor agonists. Within this previous study, tunneling spectra were generated and a single common peak among all of the agonists was determined; the IET probability density displayed behavior scaling with the efficacies of the agonists at the 5HT_{2A} receptor (36). Herein, we test the validity of this vibrational theory of protein activation by examining the assumed active peak—as determined in the above-discussed previous work (36)—at roughly 1,500–1,650 cm^{-1} ; this energy range is also in agreement with a characteristic infrared peak for histidine receptors as determined by Chee and June (33), Chee et al. (34), and Oh (35). We have calculated the effects on the tunneling probability of several deuterium variants for a pair of well-characterized 5-HT_{2A} agonists: (*R*)-DOI and DAM-57. The structure of (*R*)-DOI can be found in Fig. 1A, hereafter referred to as structure 1; note that the locations of the candidate deuterium exchanges are highlighted for ease of discussion. Similarly, the structure of DAM-57, with its deuterium exchanges can be found in Fig. 1B and is hereafter referred to as structure 2. We have used the computational methods discussed within our previous work (36), as first described by Turin (11, 12).

Several deuteration schemes of structure 1 were considered to determine the scheme capable of maximizing the possible variation in efficacy at the receptor to a high enough degree that it is unlikely to be solely attributable to the pedestrian kinetic isotope effects. By deuteration of the ethyl sidechain of DOI we found that no such deuterium analogue would present sufficient change in the agonist efficacy. The effects of deuteration on the tunneling spectrum can be seen in Fig. 2. Within each plot in Fig. 2, the blue curve denotes the natural abundance compound and is given as a comparison. As can be seen in Fig. 2, no deuteration scheme of the alkyl sidechain produces a large effect on the tunneling spectra in the target energy region. The alteration in the tunneling probability, and thus the tunneling current, associated with these schemes is a decrease of roughly 10%. Although this is a substantial difference in terms of efficacy, anything smaller than 10% may not be convincing, as it could be attributable to the normal causes of kinetic isotope effects. Additionally, alteration of isotopes changes the vibrations, possibly highlighting a kinetic isotope effect that may be 8–10% itself (85, 86), leading to a further disregard of these findings.

Experimental Procedures

In Fig. 3 we have plotted the probability distribution function (PDF) for the IET during the subsequent deuteration of one (d_3), and then both (d_6) methoxy groups on the aromatic ring of DOI. Within Fig. 3, Pro-protium DOI is shown in blue, d_3 in orange, and d_6 in green. Notably, d_6 results in an $\sim 50\%$ depletion in the assumed active peak. Such a difference should be experimentally evident; additionally this effect would be larger than a typical kinetically derived isotope effect (85, 86). Reducing the tunneling probability would result in a much lower probability of an electron completing a tunneling transfer assisted by vibrational modes within this energy range. This depletion of the associated tunneling probability in a specific energy region is attributable to two interrelated effects: (i) Isotope exchange results in shifting of energy quanta of a specific vibrational mode. (ii) Alteration of coupling between a vibrational mode (oscillating dipole) and an electron is dependent on the size of the dipole (displacement), and the relative

Table 1. Binding affinities of (*R*)-DOI and its hexadeutero isotopologue at cloned human receptors

Ligand	h5-HT _{2A} pKi (Ki)*	h5-HT _{2B} pKi (Ki)	h5-HT _{2C} pKi (Ki)
<i>R</i> -DOI HCl	8.19 ± 0.09 (6.45)	8.71 ± 0.07 (1.95)	7.96 ± 0.06 (10.96)
<i>R</i> -d ₆ -DOI HCl	8.02 ± 0.07 (9.52)	8.65 ± 0.07 (2.24)	7.99 ± 0.06 (10.23)

Values were determined by PDSP; $n = 2$ full displacement curves except for 5-HT_{2A}, where (*R*)-DOI $n = 6$ and (*R*)-d₆-DOI $n = 7$.

* (Ki) values are expressed as nanomolar.

Table 2. Comparison of (*R*)-DOI and DAM-57 and their hexadeutero isotopologues for activation of 5-HT_{2A/2B/2C} measuring Gq-mediated calcium flux, as illustrated in Fig. 4

Ligands	Gq calcium flux					
	5-HT _{2A}		5-HT _{2B}		5-HT _{2C}	
	EC ₅₀ , nM (pEC ₅₀ ± SEM)	E _{max} % 5-HT	EC ₅₀ , nM (pEC ₅₀ ± SEM)	E _{max} % 5-HT	EC ₅₀ , nM (pEC ₅₀ ± SEM)	E _{max} % 5-HT
5-HT	0.83 (9.08 ± 0.02)	100	1.29 (9.01 ± 0.12)	100	0.25 (9.62 ± 0.10)	100
(<i>R</i>)-DOI	0.58 (9.25 ± 0.08)	95 ± 4	4.80 (8.43 ± 0.12)	94 ± 1	2.19 (8.70 ± 0.14)	101 ± 2
(<i>R</i>)-d ₆ -DOI	0.63 (9.22 ± 0.10)	96 ± 4	4.40 (8.45 ± 0.10)	91 ± 1	2.23 (8.70 ± 0.15)	100 ± 3
DAM-57	1.54 (8.82 ± 0.06)	98 ± 1	13.6 (7.87 ± 0.04)	73 ± 2	57.5 (7.24 ± 0.02)	81 ± 6
d ₆ -DAM-57	1.51 (8.84 ± 0.07)	98 ± 1	12.6 (7.91 ± 0.05)	74 ± 1	49.2 (7.31 ± 0.02)	83 ± 6

Calcium flux data were acquired with human 5-HT_{2A}, 5-HT_{2B}, and 5-HT_{2C} INI-expressing tetracycline-inducible HEK cells. Estimates of EC₅₀ and E_{max} represent the average and SE of the mean (SEM) from three independent experiments performed in triplicate. E_{max} is defined as percentage of 5-HT maximum response.

magnitude of an IET PDF peak associated with a mode is dependent on the mass of an atom. This depletion of electron transfer should result in fewer successful activations of the protein than the natural abundance compound. It is for these reasons that the deuteration schemes involving exchange of all six of the methoxy hydrogens of DOI and the methyl chains on the amide of DAM-57 were selected as candidates.

d₆-(*R*)-DOI was prepared by an asymmetric synthesis, as shown in Fig. S1. NMR, mass spectral, and melting-point data were consistent with a previously published synthesis of the protio compound (87). d₆-DAM-57 was prepared by a standard route (Fig. S2), using either dimethylamine or d₆-dimethylamine. First, DOI and d₆-DOI were submitted to the National Institute of Mental Health (NIMH)-sponsored Psychoactive Drug Screening Program (PDSP) (88) to determine their binding affinities at the human 5-HT_{2A} receptor and then all compounds were tested at 5-HT_{2A}, 5-HT_{2B}, and 5-HT_{2C} receptors, measuring Gq-mediated calcium flux. A complete discussion of the synthetic routes for all molecules, as well as a discussion of the biological assays, is given in [Supporting Information](#).

Results and Discussion

Herein, we operate under the following null hypothesis: A vibration-sensitive mechanism is shared between all members of the GPCR class of proteins, possibly due to conserved topographical structures even without conservation of sequence identities, implying that general topographical structures of the GPCRs are essential in performing their biological task (89). Following this logic, it seems reasonable that the family of GPCRs likely shares a common fundamental activation mechanism. Furthermore, forms of both the lock-and-key and the glove-and-hand models exist in olfactory research, admitting acceptance of common aspects. Due to the likelihood of a common fundamental activation mechanism, findings at CNS GPCRs will have implications at mammalian ORs. We have used a series of well-characterized GPCRs (explicitly h5-HT_{2A}, h5-HT_{2B}, and h5-HT_{2C}) with two established ligands: (*R*)-DOI and DAM-57.

IET spectra were generated for both molecules, using methods prescribed by Turin in previous works (11, 12). Within a previous work (36), we examined the IET spectrum of several agonists and determined a consistently shared peak between the agonist's spectra, the area of this shared spectral aspect roughly scaled with the agonist's efficacy. This peak was consistent with the works of Chee and June (33), Chee et al. (34), and Oh (35), who attempted to generate a novel pharmacophore tool based on shared vibrational peaks. From the candidate deuteration schemes discussed above, we sought the greatest possible alteration in efficacy at the previously suggested activation peak to focus our experimental effort (36).

Both (*R*)-DOI and (*R*)-d₆-DOI were synthesized and were assayed for binding affinity (Table 1). All four compounds were assayed for functional activity, using calcium flux assays (Fig. 4 and Table 2) at the 5-HT_{2A}, 5-HT_{2B}, and 5-HT_{2C} receptors. Neither (*R*)-d₆-DOI nor d₆-DAM-57 presented significant alterations in either binding or potency and efficacy (as noted by EC₅₀ and E_{max}, respectively) compared with the protio versions. The endogenous agonist 5-HT was included in the calcium flux assays for comparison.

The process of protein agonism/antagonism involves, at minimum, two steps: binding of the ligand to the active site of the protein and the activation of the protein. It should be noted that these actions may happen in concert as proposed in the hand-in-glove/multiconformation (5–8) models or as two individual steps (11, 60, 67, 80, 90). In Table 1, we present the results of binding displacement assays comparing the relative binding affinity of both (*R*)-DOI and (*R*)-d₆-DOI at several serotonin GPCRs. Alteration in the binding kinetics, as defined by pK_i, shows no significant difference between the protonated and deuterated variants; explicitly, any difference found in the power-scaled equilibrium constant exists within the standard errors of the number, and therefore no claim of a difference can be made with confidence. The kinetic isotope effect may also affect the binding kinetics of the G protein and thus could appear independently of ligand-binding effects but would be apparent in activation studies.

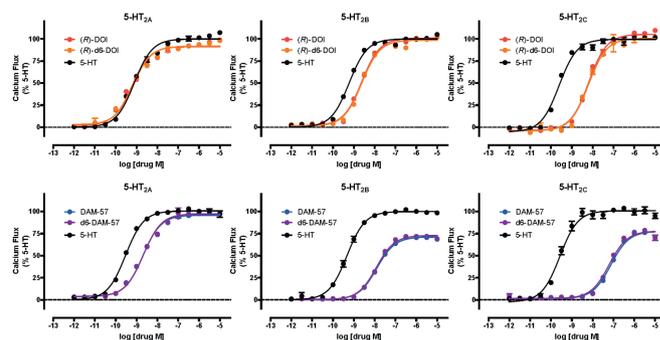


Fig. 4. Calcium flux responses at human 5-HT_{2A}, 5-HT_{2B}, and 5-HT_{2C} INI receptors for (*R*)-DOI (red), (*R*)-d₆-DOI (orange), DAM-57 (blue), and d₆-DAM-57 (purple). Data are representative at each receptor type performed in triplicate and in parallel.

Activation analysis was conducted through a series of experiments at human tetracycline-inducible HEK cells expressing all members of the h5-HT₂ receptor subclass. Receptor activation was determined by calcium flux assay, dependent on G α q dissociation. The experiment was performed in triplicate ($n = 3$) to achieve relevant statistics. Results of the G α q flux can be found in both Fig. 4 and Table 2 for five species. It is clear from Fig. 4 that DOI is a near full agonist at all three 5-HT₂ receptors examined in this assay, whereas DAM-57 shows partial agonist activity at 5-HT_{2B} and 5-HT_{2C} receptors. Additionally—and more prescient—there is no significant difference in calcium flux between either pair of parent–isotopologue compounds at any of the 5-HT₂ receptors. This conclusion is illustrated in Table 2, which gives both the EC₅₀ and the percentage of E_{max}. The power-scaled pEC₅₀, which estimates differences in potency between each pair of parent–isotopologue compounds, is within the SE and likewise for the percentage of maximum response (where 100% is taken to be the response to the endogenous 5-HT ligand). Therefore, no significant ligand-binding or activation (due to either ligand–receptor or ligand–receptor–G protein interactions) effects were found.

Herein, we found no exceptional dependency on isotopic exchange for the activity of either (R)-DOI or DAM-57. These findings do not argue for the irrelevancy of isotopic exchange, but do relegate the majority of this concern to binding. Isotope effects on the binding of a ligand with a protein have been examined extensively (85, 86) and were the subject of a recent paper on the H₂ histamine receptor (58). Whereas Kržan et al. (58) detected small alterations in the binding of ligand to receptor due to isotope exchange and conclude with possibly relevant comments concerning the VTO, we considered both the affinity and the functional activity. By considering both aspects, we are able to draw conclusions regarding the activation of the protein—addressing the VTO directly—while retaining knowledge of possible binding effects. As this work was to evaluate the dependency of activation on vibrational modes of ligands in GPCRs within the CNS through an IET-esque mechanism, we state that

we found no evidence suggestive of the plausibility of said theory. We believe that the findings within this paper argue very strongly against the VTO. Furthermore, this places our work in agreement with the transition state theories (recent works: refs. 5–8). Physicochemical properties of the ligand—other than vibrational quanta—are likely involved in the activation of both ORs—as suggested for *Drosophila* receptors by Saberi and Seyed-allaei (91)—and GPCRs in general.

Conclusion

Herein, we attempted to examine the viability of a contentious theory of protein activation—originally proposed for ORs—with a series of serotonin receptors widely expressed in the CNS: h5-HT_{2A}, h5-HT_{2B}, and h5-HT_{2C}. We have tested two well-characterized ligands—and specific isotopologues—at the 5-HT receptor family: (R)-DOI and DAM-57. Our calculations of the tunneling probability—based on Turin’s theory—predicted as much as a 50% loss of potency for both (R)-d₆-DOI and d₆-DAM-57, compared with their protium counterparts. The minor deviations in the binding between (R)-DOI and (R)-d₆-DOI were within the SE and therefore cannot be assumed to be of any significance, as were the changes in receptor function measured by calcium flux. Similarly, no deviation in the potency or efficacy was detected for d₆-DAM-57. As all values were within the SE, we can only conclude that alteration of the hydrogen isotopes at (R)-DOI and DAM-57 agonists has no significant effect on the activity at the series of 5-HT receptors studied herein. Clearly, our results are not consistent with predictions made under the VTO acting within nonolfactory GPCRs. This calls into question the viability of the IET mechanism within non-OR GPCRs, while additionally making it more difficult to argue in favor of the VTO without invoking exceptionalism of ORs within the GPCR class of proteins.

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