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### Design, Synthesis and Evaluation of Potent and Selective Ligands for the Dopamine 3 (D<sub>3</sub>) Receptor with a Novel *in vivo* Behavioral Profile

Jianyong Chen<sup>1</sup>, Gregory T. Collins<sup>2</sup>, Jian Zhang<sup>1</sup>, Chao-Yie Yang<sup>1</sup>, Beth Levant<sup>4</sup>, James Woods<sup>2</sup>, and Shaomeng Wang<sup>1,2,3,\*</sup>

1Department of Internal Medicine, University of Michigan, 1500 East Medical Center Drive, Ann Arbor, Michigan 48109

**2**Department of Pharmacology, University of Michigan, 1500 East Medical Center Drive, Ann Arbor, Michigan 48109

**3**Department of Medicinal Chemistry, University of Michigan, 1500 East Medical Center Drive, Ann Arbor, Michigan 48109

4Department of Pharmacology, Toxicology, and Therapeutics, University of Kansas Medical Center, Kansas City, Kansas 66160

#### Abstract

A series of compounds structurally related to pramipexole were designed, synthesized and evaluated as ligands for the dopamine 3 (D<sub>3</sub>) receptor. Compound **12** has a K<sub>i</sub> value of 0.41 nM to D<sub>3</sub> and a selectivity of >30,000- and 800-fold over the D<sub>1</sub>-like and D<sub>2</sub> receptors, respectively. Our *in vivo* functional assays showed that this compound is a partial agonist at the D<sub>3</sub> receptor with no detectable activity at the D<sub>2</sub> receptor.

Dopaminergic neurotransmission is mediated by five dopamine receptors  $(D_1-D_5)$ , which can be grouped into the  $D_1$ -like  $(D_1 \text{ and } D_5)$  and  $D_2$ -like  $(D_2, D_3 \text{ and } D_4)$  receptor subtypes. Recent studies have suggested that the  $D_3$  receptor is a promising therapeutic target for a variety of conditions, including drug abuse, restless legs syndrome, schizophrenia, Parkinson's disease, and depression.<sup>1-6</sup> Considerable effort has been devoted in recent years to the discovery and development of potent and selective  $D_3$  ligands.<sup>6-22</sup>

Despite intense research efforts, design of truly selective  $D_3$  ligands with good solubility and bioavailability remains a challenge. Compound 1 (pramipexole) is a potent  $D_3$ -preferring agonist but has limited selectivity over the  $D_2$  receptor *in vitro*<sup>23</sup> and *in vivo*.<sup>24,25</sup> Compound 2 was initially reported as a  $D_3$  partial agonist and has a 67-fold selectivity over the  $D_2$  receptor. <sup>2</sup> A number of potent and selective  $D_3$  ligands, such as 3, have been designed based upon the core structure of 2.<sup>17</sup> Our laboratory has reported the design of 4 as a potent and selective  $D_3$  ligand using the hexahydropyrazinoquinoline as the core structure.<sup>21</sup> Despite its relatively high affinity and excellent selectivity for  $D_3$  over other dopamine receptor subtypes, 4 has a poor aqueous solubility, which limits its *in vivo* evaluations. The poor aqueous solubility is also a major limitation for many recently described potent and selective  $D_3$  ligands and an obstacle for evaluation of these novel agents in behavioral models in animals and their therapeutic potential.

To whom correspondence should be addressed. Phone: 734-615-0362. Fax: 734-647-9647. E-mail: shaomeng@umich.edu.

To overcome this major limitation, we investigated other core structures for the design of potent and selective  $D_3$  ligands. Among them, the core structure in **1** has a number of very attractive features. First, **1** itself is a very potent  $D_3$  ligand and has a  $K_i$  value of 0.78 nM to  $D_3$  in our binding assay (Table 1). Second and importantly, **1** has an excellent aqueous solubility. Third, pramipexole dihydrochloride has been approved for the treatment of Parkinson's disease and restless legs syndrome and has an excellent pharmacological and toxicological profile in humans and in animals. Hence, **1** represents a particularly attractive template for the design of potent and selective  $D_3$  ligands with desirable physiochemical and pharmacological properties. Of note, although **1** has been widely used as a  $D_3$  preferring ligand, it potently binds to the high affinity state of the  $D_2$  receptor with a  $K_i$  value of 3.1 nM in our binding assays (Table 1), thus displaying only a 4-fold selectivity for the  $D_3$  receptor over the  $D_2$  receptor.

Recently, the crystal structures for the human  $\beta 2$  adrenergic ( $\beta 2AD$ ) G-protein coupled receptor (GPCR) were solved.<sup>26,27</sup> We have modeled the human D<sub>3</sub> receptor structure based upon the high-resolution crystal structures of  $\beta 2AD$  receptor since these two proteins belong to the same GPCR sub-family <sup>28</sup> and share close sequence homology. Because the crystal structure of  $\beta 2AD$  receptor was solved with an inverse agonist bound to it, our modeled D<sub>3</sub> structure likely represents the conformational state bound to either antagonists or inverse agonists. Hence, care must be taken when using the structure to model the interactions of the D<sub>3</sub> receptor with its ligands with different intrinsic functions. Nevertheless, we reasoned the modeled human D<sub>3</sub> structure based upon the very first human GPCR structure could be useful to guide the design of novel D<sub>3</sub> ligands.

To this end, we modeled the binding of **1** to the  $D_3$  receptor structure through computational docking, followed by extensive refinement (Supporting Information). The predicted model (Figure 3) showed that the primary amino group in the thiazol ring of **1** forms a hydrogen bonding network with the hydroxyl groups of Ser192 and Ser193. The thiazol ring in **1** is parallel to the imidazole ring in His349, making favorable  $\pi$ - $\pi$  stacking interaction. The protonated nitrogen in **1** forms a salt bridge with the negatively charged Asp110. The n-propyl group in **1** inserts into a hydrophobic channel formed by Cys114, Phe345, Phe346, Trp342 and Try373.

The predicted model of **1** in complex with the  $D_3$  receptor suggested that there is ample room available to accommodate a much larger hydrophobic group where the n-propyl group in **1** binds. Interestingly, in the adjacent area, there is another well-defined but smaller hydrophobic cavity formed by Cys114, Phe197 and Trp342 residues. We have thus designed and synthesized compound **5** to explore the interactions with these two pockets.

Compound **5** was tested for its binding affinities to the dopamine receptors using the same methods as described previously (Table 1).<sup>21</sup> It was found that **5** has a  $K_i$  value of 0.043 nM to the  $D_3$  receptor, being 18-times more potent than **1**. Compound **5**, however, also potently binds to the high affinity state of the  $D_2$  receptor with a  $K_i$  value of 2.7 nM, thus displaying a 62-fold selectivity for the  $D_3$  receptor over the  $D_2$  receptor. Similar to **1**, **5** has a weak affinity to the  $D_1$ -like receptor, its selectivity over the  $D_2$  receptor is modest.

In our previous design of **4**, we have shown that introduction of a *trans*-cyclohexyl group into the linker region yielded new ligands with much improved selectivity for the  $D_3$  receptor over the  $D_2$  receptor as compared to a linear 4-carbon linker.<sup>21</sup> We have thus designed compound **6** to investigate if introduction of this rigid cyclohexyl group into **5** may also improve the selectivity. Compound **6** binds to the  $D_3$  and  $D_2$  receptors with K<sub>i</sub> values of 0.40 nM and 307 nM, respectively. Hence, **6** is a potent  $D_3$  ligand and displays an excellent selectivity of 763-fold for the  $D_3$  receptor over the  $D_2$  receptor.

We next designed and synthesized compounds **7–10** to investigate the importance of the npropyl group in **6** for binding and selectivity. Compound **7** with an n-butyl group has a slightly weaker affinity for the D<sub>3</sub> receptor than **6** and exhibited a 2-site competition curve at the D<sub>2</sub> receptor, with roughly 10-fold less selectivity for the D<sub>3</sub> receptor over the D<sub>2</sub> receptor with the high affinity binding component. Compound **8** with an isopentyl group is 5-times less potent than **6** to the D<sub>3</sub> receptor but has a similar binding affinity to the D<sub>2</sub> receptor. Compound **9** with a bulky cyclohexylethyl group is 55-times less potent than **6** to the D<sub>3</sub> receptor but is only 3-times less potent than **6** to the D<sub>2</sub> receptor. Compound **10** with a hydrogen atom at this site has a K<sub>i</sub> value of 7.6 nM to the D<sub>3</sub> receptor, being 19-times less potent than **6**, but their binding affinities to the D<sub>2</sub> receptor are essentially the same. Therefore, our binding data clearly showed that the substitution on this nitrogen atom has a major effect on the binding to the D<sub>3</sub> receptor but modest influence on the binding to the D<sub>2</sub> receptor. Our data also showed that the n-propyl group in **6** enhances the binding affinity to the D<sub>3</sub>-receptor by 19-fold as compared to a hydrogen atom in **10**.

We next investigated the influence of the naphthyl group in **6** for binding and selectivity. Compound **11**, in which the naphthyl group is replaced by a 2-benzofuran, binds to the D<sub>3</sub> receptor with the same affinity ( $K_i = 0.51$  nM) as **6** but its selectivity over the D<sub>2</sub> receptor is decreased to 133-fold, due to its increased binding affinity to the D<sub>2</sub> receptor. Compound **12**, in which a cinnamyl group is used to replace the naphthyl, retains a high binding affinity for the D<sub>3</sub>-receptor ( $K_i = 0.41$  nM) and displays 800- and >30,000-fold selectivity over the D<sub>2</sub> and D<sub>1</sub>-like receptors. These data suggested that the modifications of the naphthyl group can have a significant effect on the selectivity, and this region should be further investigated for the design of potent and selective D<sub>3</sub> ligands.

The synthesis of compounds 5–12 is provided in the Supporting Information.

Compounds **5**, **6** and **12** were found to have good aqueous solubility. For example, the dihydrochloride salt form of **6** has an aqueous solubility greater than 100 mg/ml. Their excellent aqueous solubility provided us with an opportunity to evaluate their *in vivo* functional profiles in animals.

Another challenge in the development of selective  $D_3$  ligands was that the current *in vitro* functional assays for the  $D_3$  receptor are not predictive of the *in vivo* function of  $D_3$  ligands. <sup>6</sup> Furthermore, there was also the lack of a robust *in vivo* functional assay for the  $D_3$  receptor. To addresses these challenges, we have recently validated *in vivo* functional assays for the  $D_3$  and  $D_2$  receptors.<sup>24,25</sup> Our studies showed that yawning in rats provides a sensitive measure of *in vivo* agonist activity at the dopamine  $D_3$  receptor, <sup>24,25</sup> while the inuction of hypothermia has been shown to be mediated by agonist activity at the  $D_2$  receptor.<sup>32–33</sup> Employing these well validated assays, we evaluated **5**, **6** and **12** for their *in vivo* functional activity at the  $D_3$  and  $D_2$  receptors. Compound **1**, a known  $D_3$  and  $D_2$  agonist, was used as a control in our evaluations. The results are shown in **Figure 4**.

Consistent with the data obtained in previous studies,  $^{24,25}$  increases in yawning were observed over low doses (0.01 to 0.1 mg/kg) of **1** with inhibition of yawning and the induction of hypothermia occurring at higher doses. These data indicate that **1** functions as a preferential D<sub>3</sub> agonist *in vivo* and a D<sub>2</sub> agonist at higher doses.

Compound 5 induced yawning and produced an inverted U-shaped dose-response curve. The maximum levels of yawning induced by 5 are very similar to that induced by 1. Furthermore, hypothermia was induced by 5 at higher doses, concurrent with deceases in yawning. These data showed that 5 functions as a full agonist at the  $D_3$  and  $D_2$  receptors *in vivo*, consistent with the 2-site competition curve observed in the [<sup>3</sup>H]spiperone binding assay for 1 and 5 (Supporting Information). Furthermore, the *in vivo* data suggested that 5 is bioavailable.

Unlike 1 and 5, the dose-response curves for 6 and 12 induced yawning were relatively flat, and failed to reach significance during the initial 30 min observation period. While significant levels of yawning induced by 6 and 12 were observed after 60 min, the dose-response curves for both compounds remained relatively flat. Moreover, 6 and 12 failed to induce changes in body temperature over the initial hour of observation, an effect that is indicative of  $D_2$  agonist activity. Together, the low levels of yawning, combined with the absence of any hypothermic effect suggested two possibilities: (1) 6 and 12 function as weak partial agonists at the  $D_3$  receptor, with no detectable agonist activity at the  $D_2$  receptor; or (2) they are simply not bioavailable.

To investigate these two possibilities, we next evaluated the ability of 12 to alter compound 1-induced yawning and hypothermia and the data are shown in Figure 4. Similar to the effects of 12 alone, but unlike the effects of D<sub>3</sub>-selecitve antagonists, 24,25 low levels of yawning were observed during the initial 30 min after administration of either 10.0 or 32.0 mg/kg of 12. Interestingly, this effect appeared to persist upon administration of low doses of 1 as significant increases in yawning were observed when rats were pretreated with 12 (10.0 or 32.0 mg/kg). However, 12 resulted in a dose-dependent decrease in the amount of yawning observed following the maximally effective dose of 1 at 0.1 mg/kg. No significant effects of 12 were observed at higher doses of 1 (0.32 and 1 mg/kg). These data suggested that 12 is capable of antagonizing the D<sub>3</sub>-mediated effects of 1. However, the profile of activity for 12 is different from that observed for selective D<sub>3</sub> antagonists, which generally produce selective rightward and/or downward shifts of the ascending limb of the yawning dose-response curve for D<sub>3</sub>preferring agonists without increasing the amount of yawning observed at low doses.<sup>24,25</sup> In fact, the effects of 12 alone, and in combination with 1, suggest that it is more similar to the partial agonist, aripiprazole,<sup>34</sup> than an antagonist. Moreover, **12** failed to alter the induction of hypothermia by 1, an effect that is indicative of  $D_2$  agonist activity, which can be reliably blocked by both selective and non-selective  $D_2$  antagonists.<sup>32,33</sup> Together, our data provide evidence that 12 is a partial agonist at the D<sub>3</sub> receptor with no detectable agonist or antagonist activity at the D<sub>2</sub> receptor, thus possessing a novel in vivo functional profile.

In summary, a series of enantiomerically pure pramipexole derivatives have been designed, synthesized, and evaluated for their binding and selectivity to the  $D_3$ ,  $D_1$ -like and  $D_2$  receptor. This led to the identification of several potent and highly selective  $D_3$  ligands with excellent aqueous solubility. Our *in vivo* functional evaluations showed that while **5** functions as a full  $D_3$  agonist, **12** behaves as a selective  $D_3$  partial agonist with no activity at the  $D_2$  receptor. Further *in vivo* studies are underway to evaluate the therapeutic potential of **12** for the treatment of drug abuse and other indications. The results will be reported in due course.

#### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

#### Abbreviations

 $D_{1-5}$ , dopamine 1–5 receptor subtypes;  $\beta$ 2AD, the human  $\beta$ 2 adrenergic receptor; GPCR, G-protein coupled receptor.

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**Figure 1.** Chemical structures of representative D<sub>3</sub> ligands.



**Figure 2.** New analogues structurally related to pramipexole.

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#### Figure 3.

Predicted binding model of compound 1 to the human  $D_3$  receptor. For protein, carbon atoms of human  $D_3$  are shown in white, oxygen atoms in red, and nitrogen atoms in blue. Side chains of crucial residues in the binding site are shown as stick and labeled. Hydrogen bonds between 1 and  $D_3$  are depicted in dotted line in yellow. Figures were generated by Pymol.

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#### Figure 4.

Functional evaluations of the  $D_3$  and  $D_2$  activity of pramipexole, compounds **5**, **6** and **12** in yawning and hypothermia assays in rats. Top and middle panels: Induction of yawning or hypothermia by  $D_3$  ligands. Bottom panels: Interactions between pramipexole and compound **12** in yawning and hypothermia assays.

# Table 1

Binding affinities at the  $D_1$ -like,  $D_2$  and  $D_3$  receptors in binding assays using rat brain. Data represent the mean  $\pm$  SEM of 3–5 independent determinations. For compounds producing a 2-site fit in competition with [<sup>3</sup>H]-spiperone, K<sub>i</sub> values are presented for the high and low affinity components and are indicated by the designation "(h)" or "(l)". All other K<sub>i</sub> values are based on a single-site model.

		$\mathbf{K}_{\mathbf{i}} \pm \mathbf{SEM} \ (\mathbf{nM})$		Sel	ectivity
Ligand	$D_{3}[^{3}H]PD128 907$	D <sub>2</sub> [ <sup>3</sup> H]Spiperone	D <sub>1</sub> -like [ <sup>3</sup> H]SCH23390	D <sub>2</sub> –like /D <sub>3</sub>	D <sub>1</sub> –like /D <sub>3</sub>
1	0.78	$3.1 \pm 0.3$ (h) $6400 \pm 1700$ (l)	>100,000	4.0	>100,000
4	$5.7\pm0.4$	>10000	>50000	>1000	>5000
5	$0.043\pm0.006$	$2.7 \pm 0.4$ (h) $6700 \pm 1500$ (l)	$11,000\pm500$	62	>100,000
9	$0.40\pm0.057$	$307 \pm 38$	$3,400\pm300$	763	>7,000
7	$0.74\pm0.083$	$55 \pm 12$ (h) $1300 \pm 180$ (l)	$5,400\pm500$	74	>7,000
8	$2.2\pm0.10$	$345 \pm 33$	$13,000 \pm 1,000$	157	>5,000
6	$23 \pm 2.7$	$1,200\pm170$	$4,400\pm800$	53	194
10	$7.6\pm0.87$	$670 \pm 140$	$64,000 \pm 7,000$	88	>8,000
11	$0.51 \pm 0.10$	$68 \pm 4.6$	$4,900 \pm 600$	133	>9,000
12	$0.41\pm0.031$	$330 \pm 69$	$13,000 \pm 1,700$	800	>30,000