

COMPREHENSIVE PHARMACOLOGICAL PROFILING OF THE HUMAN 5-HT_{7A} RECEPTOR ISOFORM EXPOSES NOVEL SIGNALING Claire NORMAND¹, Guilhem DUGAST¹, Laurent SABBAGH¹, Thomas RAY², Paul WREN², Arturo MANCINI¹ ¹Domain Therapeutics NA Inc., Montreal, Québec, Canada ²Mindstate Design Labs, South San Francisco, CA, USA

INTRODUCTION

The 5-HT (serotonin) receptor type 7 (5-HT₇R) is one of the most recently identified members of the 5-HT receptor family. Pharmacological modulation of this G protein coupled receptor (GPCR) is considered a promising approach for the treatment of various neurological and psychiatric disorders including anxiety, depression, schizophrenia and Alzheimer's disease¹.

Human (h) 5-HT₇R has been reported to couple primarily to G α s, which activates adenylate cyclases and leads to the production of cAMP². However, it has now become evident that many GPCRs can couple to more than one signaling pathway, and that different ligands acting at a given receptor can selectively promote the activation of different subsets of these pathways. This so-called "functional selectivity" (or biased signaling) highlights the importance of exhaustively testing multiple ligands on multiple signaling pathways when defining a receptor's signaling repertoire. To date, h5-HT₇R's full signaling signature in response to different ligands remains unexplored.

In this study, we applied Domain Therapeutics' bioSens-All[®] platform of enhanced bystander Bioluminescence Resonance Energy Transfer (ebBRET)-based biosensors to profile the signaling and pharmacology of ten diverse 5-HT₇R ligands at the recombinantly expressed h5-HT_{7a}R isoform in HEK293 cells. The agonist and antagonist activity of each ligand was characterized on an extensive panel of sixteen pathway-specific biosensors. The results obtained confirmed h5-HT_{7a}R's coupling to Gas and are in alignment with previously published data³⁻⁷. Moreover, our data exposed novel couplings to brain-enriched Gi/o-family G proteins (i.e., GaoA, GaoB, $G\alpha z$), $G\alpha 13$, $G\alpha 14$ and (especially) $G\alpha 15$. Interestingly, comparison of the ligand activity on Gas and the newly identified Ga15 pathway enabled the broad classification of compounds into five functionally distinct pharmacological clusters. Among the identified clusters, one defined by LP-44 and LSD for example produced a switch in h5-HT_{7a}R G protein coupling. Specifically, ligands in this cluster displayed no detectable agonist activity on $G\alpha s$ but fully antagonized 5-HT-induced $G\alpha s$ signaling while activating $G\alpha 15$ and other pathways.

MATERIALS AND METHODS



Figure 1: Assay principle underlying our ebBRET-based biosensors⁸. G protein-specific *Renilla* luciferase (RlucII)-tagged (R) effector proteins are recruited to activated (GTP-bound) untagged G alpha subunits following activation of untagged receptor. This event brings Rlucll in close proximity to the plasma membraneanchored *Renilla* green fluorescent protein (G), leading to an increase in ebBRET. Exceptionally for the G α s biosensor, RlucII is directly fused to the Gas protein. GPCR-mediated Gas activation leads to its dissociation from the plasma membrane, resulting in a reduction in ebBRET.



Figure 2. Multiparametric profiling of tool compound pharmacology at the h5-HT_{7a}R reveals G protein coupling diversity and novel signaling via Ga15. HEK293 cells were co-transfected with pathway-specific biosensor-coding plasmids and a plasmid encoding for h5-HT_{7a}R. Ligands were assessed for both agonist and antagonist (at an EC₉₀ of 5-HT) activity. Curves were fitted using the log(agonist) vs. response - Variable slope (four parameters) non-linear regression model (GraphPad 10). Data are presented as mean values ± SEM, n=3-4. *In the absence of a clear quantifiable (EC₅₀ value) concentration response, % activity was calculated using the highest ligand concentration on the ligand's dose response curve. ND: no response or not determinable. h5-HT_{7a} exhibits a wide G protein signaling repertoire, coupling to at least one member from each of the four G protein families.

CONCLUSIONS

>h5-HT_{7a}R can signal through various G protein pathways in addition to the Gas pathway, potentially influencing a wide range of cellular processes and physiological functions. > Novel coupling through Ga15 was identified; tool compound activity on this pathway allows for their classification into five functionally distinct pharmacological clusters.

REFERENCES: 1. Fukuyama *et al.* 2023. Int J Mol Sci. Jan 20;24(3):2070; **2.** Guseva *et al.* 2014. Front Behav Neurosci. Oct 1:8:306; **3.** Krobert *et al.* 2004. J Med Chem. Dec 16;47(26):6616-245; **5.** Atanes *et al.* 2013. Pharmacol Res Perspect. Dec;1(2):e00013; **6.** Brenchat *et al.* 2009. Pain. Feb;141(3):239-247; **7.** Deau et al. 2015. J Med Chem. Oct 22;58(20):8066-96; **8.** Avet, Mancini et al. 2022. Elife. Mar 18:11:e74101.

> The data and methods reported herein enable deeper pharmacological characterization and understanding of 5-HT_{7a}R function and may help inform its therapeutic exploitation.

RESULTS

AGONIST activity										
5-HT	5-CT	8-OH-DPAT	Methiothepin	AS19	LP211	SB 269970	LP44	Serodolin	LSD	
100.00	97.84	79.28	ND	21.79	ND	ND	ND	ND	ND	
1.57	3.25	11.75	ND	3.47	ND	ND	ND	ND	ND	
5 36	1 17	1828.00		17 54			ΝП			
0.67	0.27	1058.00		13.83						
0.01	0.21	1000.00		10.00	ND	ND	ND	ND	ND	
400.00	404.00	00.04	70.04	07 70	ND	44.00	40.00	66.07	99 59	
100.00	2.95	96.01	-/ b.31	97.70		-41.08	40.92	-00.U/ 2.50	82.32	
5.20	3.05	4.70	5.21	5.21	ND	2.21	5.05	2.59	4.70	
0.30	0.23	19.04	9.63	1.51	ND	5.13	10.86	2.52	3.00	
0.07	0.05	4.90	5.07	0.48	ND	2.57	4.62	0.80	0.96	
100.00	77.53	31.53	ND	15.54	ND	ND	ND	ND	ND	
3.73	3.53	4.82	ND	3.89	ND	ND	ND	ND	ND	
2.16	2.15	559.20	ND	28.37	ND	ND	ND	ND	ND	
0.53	0.38	293.30	ND	26.51	ND	ND	ND	ND	ND	
100 00	109 80	68.38	-45 22	59 24	ND	-23 93	10.83	-27 20	36 70	
4 18	4 58	3.88	8 25	4 16		1.92	2 60	6.62	3 14	
		0.00	0.20		ne -		2.00	0.02	0.11	
0.93	0.56	97.34	1067.00	4.37	ND	4.82	3.17	29.36	1.96	
0.28	0.14	31.21	820.80	2.34	ND	3.47	4.55	50.74	1.14	
400.00		400.40*		400 70*		4 4 9 . 9 9			77.00	
100.00	99.90^	102.19*	-314.20	103.78^	ND	-149.60	ND	-232.50	//.63	
10.99	5.18"	3.02"	40.04	3.16"	ND	15.30	ND	16.89	32.77	
1.66	0.49	152.20	59.06	2.09	ND	7.53	ND	8.55	13.62	
1.43	0.32	226.40	41.98	1.89	ND	7.13	ND	4.94	24.05	
100.00	94.83	81.93	-17.55	51.41	ND	ND	ND	ND	20.80	
9.70	16.07	60.22	16.82	49.76	ND	ND	ND	ND	16.01	
1 9/	2 72	840 40	1 1 2	0.47					104 20	
1.04 1.17	2.13	2647.00	1272 00	7.05					306.70	
1.17	2.22	2077.00	1212.00	7.00					300.70	

			ΔΝΤΔ	GONI	ST act	ivity			
	5-CT	8-OH-DPAT	Methiothepin	AS19	LP211	SB 269970) LP44	Serodolin	LSD
	ND	ND	107.10	67.76	107.10	118.40	111.50	110.00	108.70
	ND	ND	4.46	6.80	3.91	5.26	6.81	4.27	4.91
	ND	ND	2.33	114.70	6.68	1.48	174.30	0.96	3.04
	ND	ND	0.62	66.41	1.54	0.55	44.36	0.27	1.05
1									
	ND	ND	167.70	ND	128.40	137.70	55.85	164.40	ND
	ND	ND	5.03	ND	14.41	5.11	8.04	4.02	ND
	ND	ND	6.33	ND	66.65	6.25	291.80	3.24	ND
	ND	ND	1.22	ND	39.55	1.84	193.20	0.71	ND
	ND	ND	109.10	54.66	92,54	92.60	72.28	92.13	80.25
	ND	ND	5.14	6.76	9.17	3.52	6.97	3.46	4.22
	ND	ND	2 17	ND	11 13	0 41	209 90	0 41	0 36
	ND	ND	0.99	ND	10.57	0.23	132.50	0.26	0.37
	ND	ND	155.70	ND	120.90	121.70	84.66	127.70	43.26
	ND	ND	11.79	ND	69.07	9.05	23.82	6.75	9.87
		ΝП	5 88	ND	43 40	2 81	161 10	3 46	6 1 1
	ND	ND	2.79	ND	141.00	2.48	260.20	1.60	12.80
	L								
	ND	ND	334.80	ND	156.26*	221.20	89.86*	310.20	ND
	ND	ND	21.03	ND	7.99*	26.93	6.75*	14.69	ND
	ND	ND	41.05	ND	ND	61.86	ND	40.27	ND
	ND	ND	14.15	ND	ND	45.13	ND	13.22	ND
	ND	ND	121.90	ND	93.89	95.02	123.00	110.80	39.75
	ND	ND	18.17	ND	21.81	15.41	451.00	10.04	9.16
	ND	ND	5.16	ND	21.24	6.74	0.56	3.68	4.33
	ND	ND	5.22	ND	27.76	8.04	22.09	2.54	9.75

Figure 3. h5-HT_{7a}R coupling to Ga15 is specific and receptor-dependent. HEK293 cells were cotransfected with the Galf biosensor and increasing amounts of either h5-HT_{7a}R coding plasmid (a) or $G\alpha 15$ protein-coding plasmid (b). The agonist activity of 5-HT and LSD was assessed and resulting curves were fitted using the log(agonist) vs. response - Variable slope (four parameters) non-linear regression model (GraphPad 10). Data are presented as mean values \pm SEM, n=2-3. Coupling to G α 15 is lost in cells not transfected with h5-HT_{7a}R (a) or G α 15 protein (b). Coupling efficacy was also h5-HT_{7a}R dose-dependent and G α 15 protein dose-dependent .







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% Gas Activity (vs. 5-HT)

Figure 4. Analysis of compound activity on Gas and Ga15 enables the classification of compounds into five pharmacological clusters: 1) 5-HT like compounds; 2) Inverse agonists; 3) Neutral antagonist; 4) Compounds with full agonist activity on $G\alpha 15$ and partial agonist activity on $G\alpha s$ and other pathways; 5) Compounds with no detectable agonist activity on $G\alpha s$ but with agonist activity on $G\alpha 15$ and other pathways, resulting in antagonism of 5-HT-induced G α s signaling with agonism (partial) on other pathways.