

Table S6. G_s-mediated cAMP accumulation assay results of D5R.

Potency (pEC₅₀) were extracted from a minimum of 3 independent assays in at least triplicate. pEC₅₀ displayed values are mean ± SEM. Delta FOB for difference in either Fold of Basal (FOB) or Delta pEC₅₀ when compared to wild-type receptor value. Average E_{max} and basal values were determined from “log(agonist) vs. response – Variable slope (four parameters) or log(inhibition) vs. response – Variable slope (four parameters)” function in Graphpad Prism 8.4 software (Graphpad Software Inc., San Diego, CA) and were divided by 10³ for display, basal values are enclosed with parentheses in the table column. Color scheme is based on the effects of mutations on relative pEC₅₀ and Fold of Basal (FOB) values with red for reduced potency/efficacy and blue for increased potency/efficacy when compared to wild-type values for each ligand. ND - not determined.

BW	D5-Dopamine						D5R-Rotigotine				
		Emax (Basal)	FOB	ΔFOB	pEC ₅₀	ΔpEC ₅₀	Emax (Basal)	FOB	ΔFOB	pEC ₅₀	ΔpEC ₅₀
	WT	18.2 ± 2.9 (1.8 ± 0.5)	13.2 ± 4.9	0	9.82 ± 0.08	0	17.5 ± 1.2 (1.7 ± 0.3)	11.8 ± 2.3	0	9.25 ± 0.12	0.00
2.61	K98A	29.5 ± 8.3 (1.3 ± 0.5)	24.9 ± 2.4	11.7	9.51 ± 0.02	-0.31	31.9 ± 0.1 (1.5 ± 0.1)	24.0 ± 2.6	10.8	9.29 ± 0.12	0.04
3.28	W116A	37.9 ± 8.9 (0.3 ± 0.1)	142.1 ± 5.9	128.9	7.02 ± 0.01	-2.8	24.0 ± 2.9 (0.3 ± 0.1)	97.3 ± 7.3	84.1	6.13 ± 0.11	-3.12
3.32	D120A	16.9 ± 5.2 (13.2 ± 4)	1.3 ± 0.1	-11.9	7.10 ± 1.19	-2.72	N.D.	N.D.	N.D.	N.D.	N.D.
3.33	I121A	30.2 ± 7.2 (0.2 ± 0.1)	131.9 ± 10.3	118.7	5.47 ± 0.13	-4.35	27.7 ± 4.1 (0.2 ± 0.1)	119.4 ± 13.0	106.2	5.48 ± 0.10	-3.77
3.36	S124A	36 ± 11.6 (0.7 ± 0.3)	61.2 ± 9.1	48	7.46 ± 0.14	-2.36	38.5 ± 9.2 (0.8 ± 0.3)	58.5 ± 7.1	45.3	7.39 ± 0.13	-1.86
3.37	T125A	18.8 ± 7.2 (1.2 ± 0.5)	15.8 ± 0.2	2.6	6.35 ± 0.10	-3.47	25.6 ± 5.1 (1.1 ± 0.3)	23.8 ± 1.7	10.6	8.43 ± 0.10	-0.82
3.40	I128A	18.6 ± 4.0 (0.1 ± 0.1)	124.6 ± 2.6	111.4	6.44 ± 0.10	-3.38	2.0 ± 0.5 (0.1 ± 0.1)	13.8 ± 2.0	0.6	6.75 ± 0.22	-2.50
ECL2	S219A	18.9 ± 5.0 (1.3 ± 0.3)	19.1 ± 8.0	5.9	9.87 ± 0.11	0.05	21.4 ± 3.4 (1.8 ± 0.4)	14.1 ± 4.6	0.9	9.49 ± 0.09	0.24
ECL2	L221A	N.D.	N.D.	N.D.	N.D.	N.D.	N.D.	N.D.	N.D.	N.D.	N.D.
5.42	S229A	22.4 ± 4.6 (2.0 ± 0.5)	15.3 ± 6.4	2.1	7.49 ± 0.11	-2.33	23.0 ± 4.0 (1.7 ± 0.6)	21.0 ± 6.4	7.8	6.37 ± 0.43	-2.88
5.43	S230A	35 ± 12.6 (0.2 ± 0.1)	178.6 ± 77.3	165.4	7.48 ± 0.14	-2.34	39.4 ± 9.2 (0.4 ± 0.1)	95.4 ± 12.5	82.2	7.37 ± 0.10	-1.88
5.46	S233A	33.5 ± 9.4 (2.4 ± 0.6)	18.8 ± 8.3	5.6	6.27 ± 0.14	-3.55	24.7 ± 5.2 (1.9 ± 0.6)	20.4 ± 6.9	7.2	9.84 ± 0.19	0.59
6.48	W309A	31.4 ± 10.1 (0.1 ± 0.1)	273.5 ± 27.0	260.3	5.73 ± 0.06	-4.09	7.0 ± 1.8 (0.1 ± 0.1)	55.1 ± 6.4	41.9	7.10 ± 0.11	-2.15
6.51	F312A	27.4 ± 7.4 (0.2 ± 0.1)	128.9 ± 13.9	115.7	6.16 ± 0.21	-3.66	29.6 ± 5.9 (0.2 ± 0.1)	164.7 ± 28.2	151.5	6.66 ± 0.15	-2.59
6.52	F313A	37.7 ± 16.6 (0.2 ± 0.1)	150.3 ± 22.4	137.1	6.64 ± 0.40	-3.18	42.0 ± 11.4 (0.2 ± 0.1)	161.2 ± 19.9	148.0	6.76 ± 0.21	-2.49
6.55	N316A	30.6 ± 15.2 (0.2 ± 0.1)	135.3 ± 21.0	122.1	6.72 ± 0.06	-3.1	34.1 ± 12.0 (0.1 ± 0.1)	247.6 ± 75.5	234.4	6.92 ± 0.06	-2.33
7.35	F341A	30.7 ± 9.5 (0.4 ± 0.1)	76.1 ± 0.4	62.9	7.96 ± 0.04	-1.86	34.1 ± 5.3 (0.4 ± 0.1)	86.9 ± 3.9	73.7	7.62 ± 0.08	-1.63
7.36	D342A	24.7 ± 8.2 (3.1 ± 0.3)	7.4 ± 1.8	-5.8	9.58 ± 0.18	-0.24	27.5 ± 7.3 (2.8 ± 0.5)	9.7 ± 2.3	-3.5	9.26 ± 0.25	0.01
7.39	V345A	20.4 ± 4.4 (3.8 ± 1.0)	5.5 ± 0.3	-7.7	9.78 ± 0.04	-0.04	20.4 ± 3.4 (3.8 ± 0.9)	5.6 ± 0.3	-7.6	9.91 ± 0.11	0.66
7.43	W349A	11.6 ± 3.3 (0.1 ± 0.1)	75.5 ± 13.9	62.3	6.92 ± 0.06	-2.9	1.6 ± 0.3 (0.2 ± 0.1)	9.2 ± 0.9	-4.0	6.17 ± 0.34	-3.08