SUPPORTING INFORMATION

Tryptamine-based derivatives as Transient Receptor Potential Melastatin type-8 (TRPM8) channels modulators

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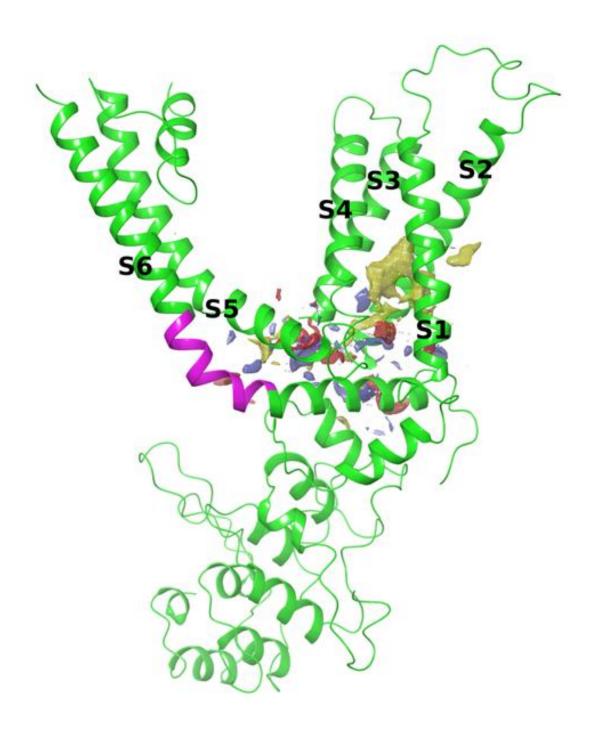


Figure S1: Sitemap graphical output for BP1. Hydrophobic surfaces are depicted in yellow, HB donors locations as blue surfaces, HB acceptors locations as red surfaces. The 980-990 region is highlighed in magenta

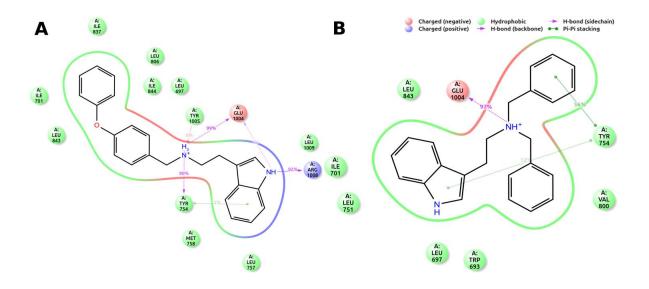


Figure S2 Protein-Ligand contacts for 21 (panel A) and 12 (panel B), recorded during 12 ns long MD trajectories.

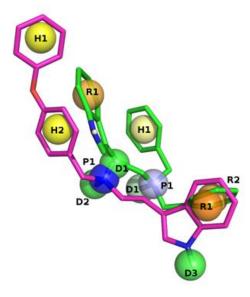


Figure S3: Pharmacophore models for tryptamine-based TRPM8 agonists and antagonists. Chemical features are depicted as spheres and color-coded as follows: yellow, H = hydrophobic; blue, P = positive charge; green, D = HB donor; orange, R = aromatic ring. Chemical features of the antagonist model are depicted in paler colors. **21** (magenta sticks) and **12** (green sticks) are represented as references.

	Agonist Model							Antagonist Model				
Comp.	H1	Н2	R1	P1	D1	D2	D3	H1	R1	R2	P1	D1
comp.	[701	I ⁸⁴⁴	Y ⁷⁵⁴ M ⁷⁵⁸	E ¹⁰⁰⁴	Y ⁷⁵⁴	E ¹⁰⁰⁴	R ¹⁰⁰⁸	Λ_{800}	Υ ⁷⁵⁴	R ¹⁰⁰⁸ Y ⁷⁵⁴	E ¹⁰⁰⁴	E ¹⁰⁰⁴
4	0	X	X	X	X	X			X	X	X	X
5		X	X	X	X	X	\		X	X	X	X
6	\	X	0	X	X	X			X	X	X	X
7		X	X	X	X	X			0	0	0	0
12			X	X		X	X	X	X	X	X	X
15		X	X	X	X	X	X		X	X	X	X
16		X	X	X	X	X	X		X	0	X	X
18		X	X	X	X	X	X		X	X	X	X
21	X	X	X	X	X	X	X		X	X	X	X
22s	0		0	X	0			X		0	X	X
22r	0		0	X		X	X	X	X	X	X	X

Table S1: Fittings of the tryptamine-based agonists and antagonists on the respective pharmacophore models. (X=good fit, O=partial fit, /=misfit, e.g. polar group on a hydrophobic location)

2. Qualitative HPLC runs for derivatives 4-12 and 14-22

