

Supporting Information

***N*-[1,3-Dialkyl(aryl)-2-oxoimidazolidin-4-ylidene]-aryl(alkyl)sulphonamides as novel selective human cannabinoid type 2 receptor (hCB2R) ligands. Insights into the mechanism of receptor activation/deactivation.**

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Figure S1. Competition radioligand binding assay of **10d** (A), **10e** (B), **10f** (C), **10g** (D), **10h** (E), **12a** (F), **12b** (G), **12e** (H) and **13** (I) at hCB2R, using [³H]CP55 940. Data are reported as mean ± SE of three independent experiments conducted in duplicate.

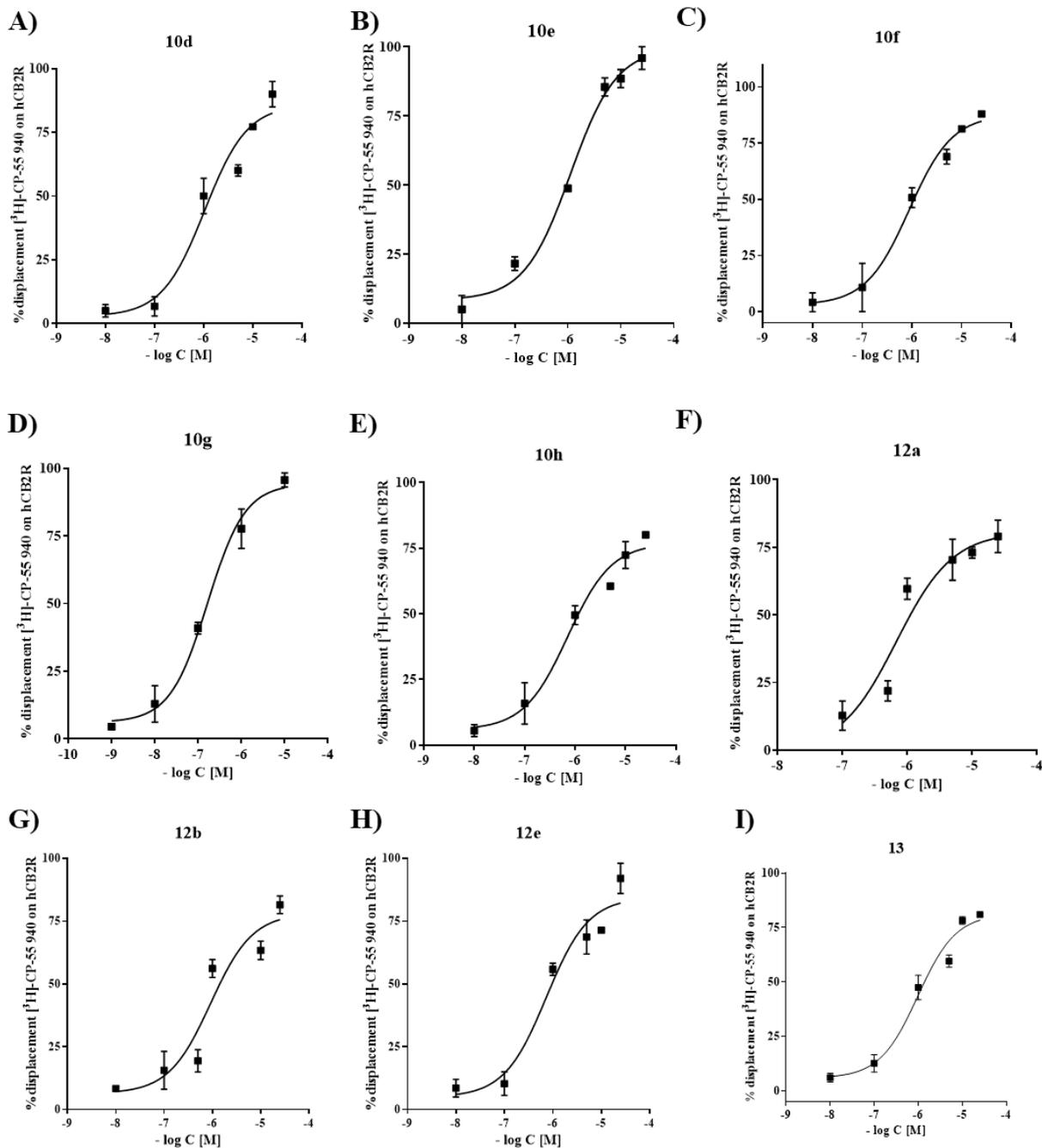


Figure S2. Chemical structure of compounds 47 (a), 49 (b), 51 (c) and 53 (d) from Mugnaini et al. [12].

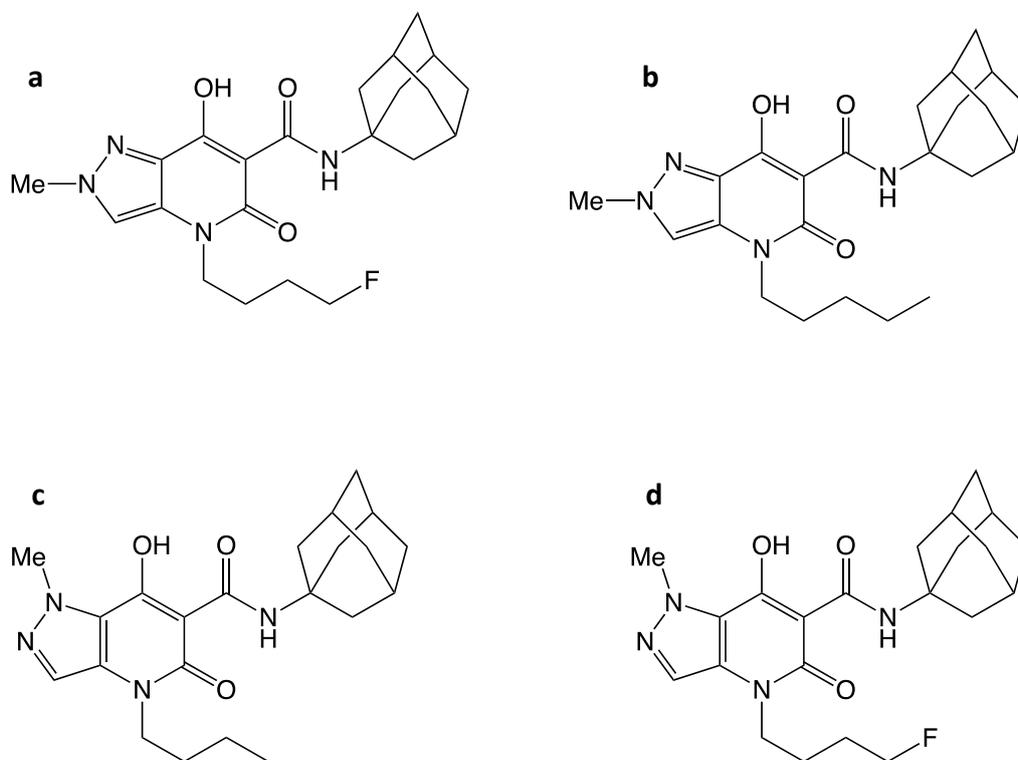


Figure S3. Docking poses of compound 49 (dark teal sticks, **a**) and compound 47 (grey sticks, **b**) in the antagonist conformation of hCB2R (PDB ID 5zty, light orange cartoon), and compound 51 (lilac sticks, **c**) and compound 53 (magenta sticks, **d**) in the agonist conformation of hCB2R (PDB ID 6kpc, light green cartoon). Compounds have been reported and described by Mugnaini et al. [12]. Residues lining the binding site are represented as capped sticks and labelled, helices are numbered as TM1-7.

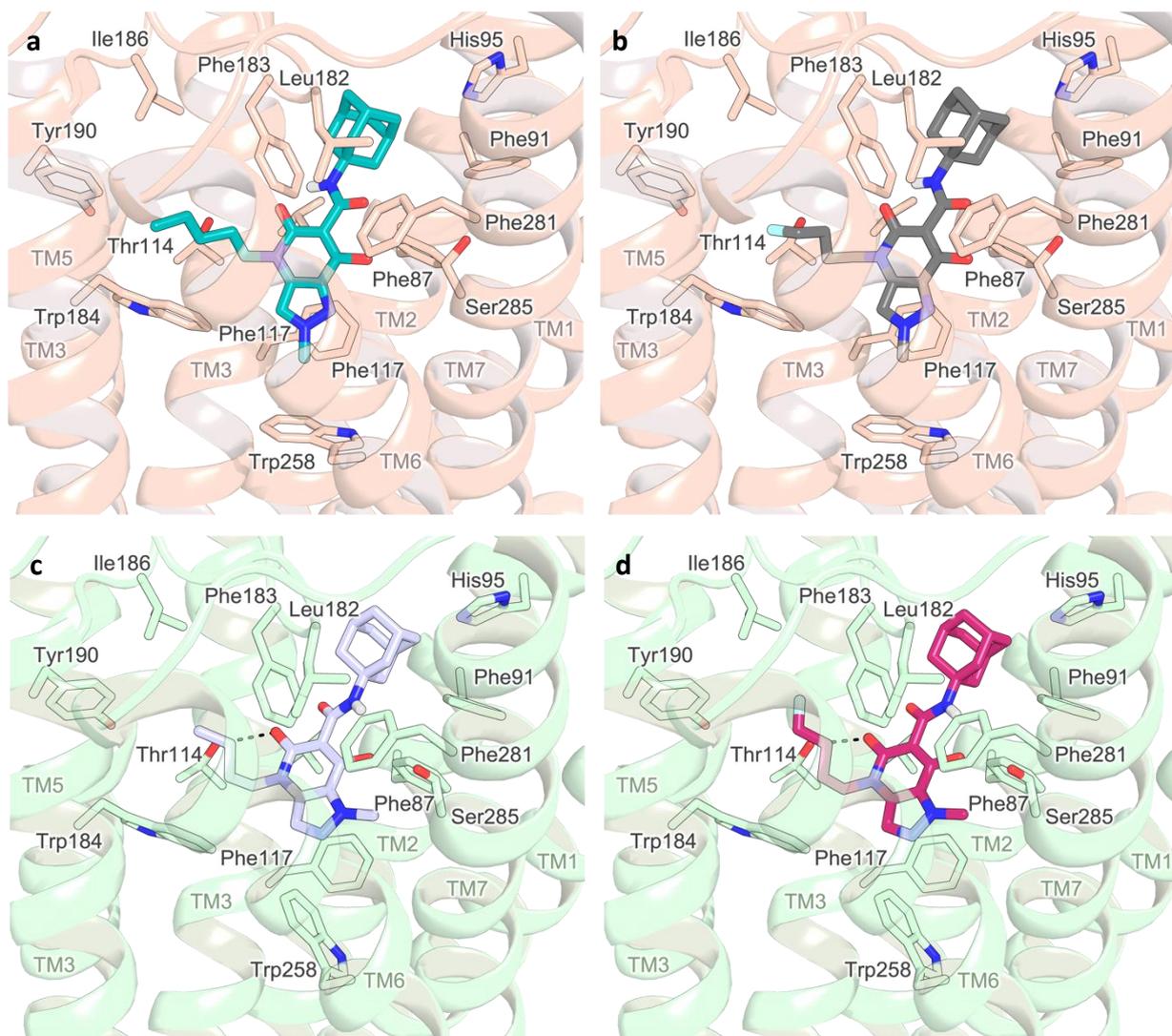


Figure S4. Docking poses of compounds **10e** (salmon pink, **a**) and **10g** (cyan, **b**) in the agonist conformation of hCB2R receptor (PDB ID: 6kpc, light green cartoon). Residues lining the binding site are represented as capped sticks and labelled, helices are numbered as TM1-7. Hydrogen bonds are represented as black dashed lines.

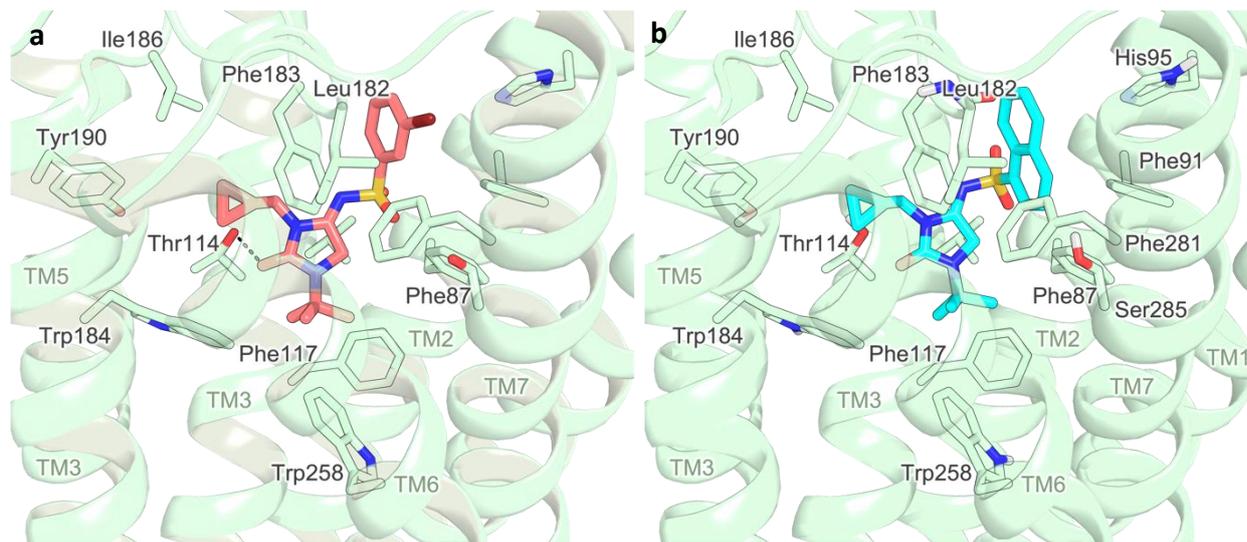


Figure S5. Docking poses of compound **12b** in the antagonist conformation of CB2R receptor (PDB ID 5zty). Residues lining the binding site are represented as capped sticks and labelled, transmembrane helices are numbered as TM1-7. Hydrogen bonds are represented as black dashed lines.

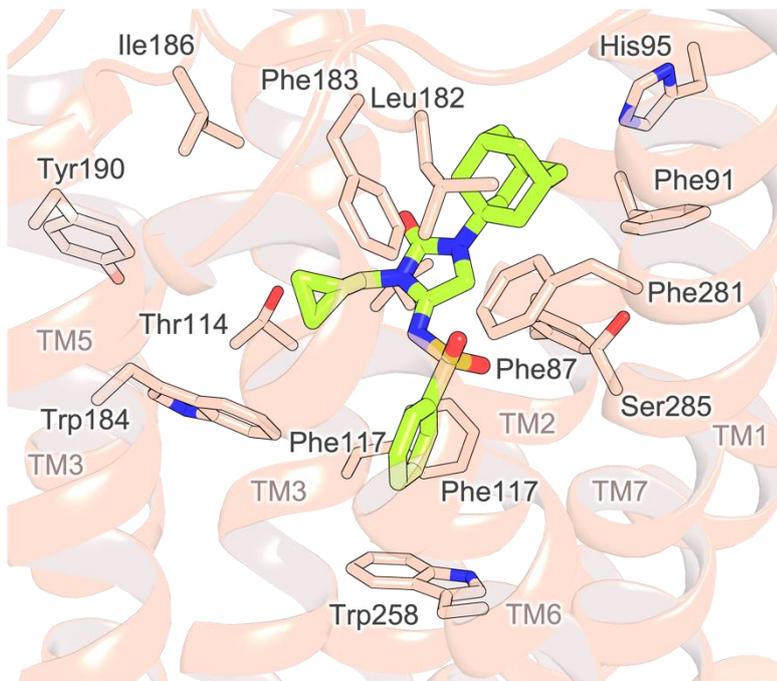


Figure S6. Alignment of **12e** docking pose (yellow) and antagonist AM10257 (green) in the antagonist-like state (PDB ID 5zty). Protein is coloured in light orange cartoon, residues lining the binding site are represented as capped sticks and labelled, helices are numbered as TM1-7.

