

Structural insights into isoquinoline small molecule ligand-induced agonism and modulation on GLP-1R

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Peptide mono- and multi-target agonists for class B1 G protein-coupled receptors (GPCRs) that co-target the glucagon-like peptide 1 receptor (GLP-1R) with one or more additional receptors are being pursued as novel therapeutics for treatment of metabolic diseases. But developing non-peptide agonists and modulators for these receptors with higher oral availability and lower manufacturing costs remains attractive for next-generation drug discovery. An isoquinoline small molecule ligand termed BI-14185 (one stereoisomer of LSN3451217) was reported to be a agonist and positive allosteric modulator (ago-PAM) at the GLP-1R^{1,2}. In the current study, we utilized cryogenic electron microscopy (cryo-EM) to understand the molecular mechanism of agonism and modulation of BI-14185 at the GLP-1R with/without the co-treatment of a peptide agonist. Two potential putative binding pockets for BI-14185 within the GLP-1R TMD were identified, one within the transmembrane helices (TM) core and the second located on the intracellular side of TM 5/6. These two binding pockets shared some similarity to binding sites identified in previously disclosed small molecule ligand-bound GLP-1R structures but they displayed unique interaction patterns with the receptor. The current study advances our understanding about BI-14185-induced agonism and modulation of peptide agonist-mediated GLP-1R responses. With future validation and investigation, this study may benefit design of novel non-peptide agonists/modulators for class B1 GPCRs.

References:

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