

# The conformational signature of $\hat{I}^2$ -arrestin2 predicts its functions

Nature

531, 665-668

DOI: [10.1038/nature17154](https://doi.org/10.1038/nature17154)

Citation Report

#	ARTICLE	IF	CITATIONS
1	Receptor antagonism/agonism can be uncoupled from pharmacoperone activity. <i>Molecular and Cellular Endocrinology</i> , 2016, 434, 176-185.	1.6	11
2	A Novel Allosteric Activator of Free Fatty Acid 2 Receptor Displays Unique Gi-functional Bias. <i>Journal of Biological Chemistry</i> , 2016, 291, 18915-18931.	1.6	66
3	Identification of key phosphorylation sites in PTH1R that determine arrestin3 binding and fine-tune receptor signaling. <i>Biochemical Journal</i> , 2016, 473, 4173-4192.	1.7	25
4	Downregulation of a GPCR by $\beta$ -Arrestin2-Mediated Switch from an Endosomal to a TGN Recycling Pathway. <i>Cell Reports</i> , 2016, 17, 2966-2978.	2.9	42
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6	Ligand-Dependent Modulation of G Protein Conformation Alters Drug Efficacy. <i>Cell</i> , 2016, 167, 739-749.e11.	13.5	113
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