## Douglas J Kojetin

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1465299/publications.pdf

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66 papers

4,537 citations

34 h-index 110387 64 g-index

80 all docs 80 docs citations

times ranked

80

6533 citing authors

#	Article	IF	CITATIONS
1	Chemical systems biology reveals mechanisms of glucocorticoid receptor signaling. Nature Chemical Biology, 2021, 17, 307-316.	8.0	11
2	CAR directs T cell adaptation to bile acids in the small intestine. Nature, 2021, 593, 147-151.	27.8	36
3	Structural mechanism underlying ligand binding and activation of PPARγ. Structure, 2021, 29, 940-950.e4.	3.3	19
4	Structural basis for heme-dependent NCoR binding to the transcriptional repressor REV-ERB $\hat{l}^2$ . Science Advances, 2021, 7, .	10.3	13
5	Assessment of NR4A Ligands That Directly Bind and Modulate the Orphan Nuclear Receptor Nurr1. Journal of Medicinal Chemistry, 2020, 63, 15639-15654.	6.4	34
6	A molecular switch regulating transcriptional repression and activation of PPAR $\hat{I}^3$ . Nature Communications, 2020, 11, 956.	12.8	45
7	Structural Basis of Altered Potency and Efficacy Displayed by a Major in Vivo Metabolite of the Antidiabetic PPARÎ <sup>3</sup> Drug Pioglitazone. Journal of Medicinal Chemistry, 2019, 62, 2008-2023.	6.4	26
8	The Tat inhibitor didehydroâ€cortistatin A suppresses SIV replication and reactivation. FASEB Journal, 2019, 33, 8280-8293.	0.5	17
9	Didehydro-Cortistatin A Inhibits HIV-1 by Specifically Binding to the Unstructured Basic Region of Tat. MBio, 2019, 10, .	4.1	56
10	Quantitative structural assessment of graded receptor agonism. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 22179-22188.	7.1	21
11	PGRMC2 is an intracellular haem chaperone critical for adipocyte function. Nature, 2019, 576, 138-142.	27.8	96
12	Defining a Canonical Ligand-Binding Pocket in the Orphan Nuclear Receptor Nurr1. Structure, 2019, 27, 66-77.e5.	3.3	37
13	Cryptic glucocorticoid receptor-binding sites pervade genomic NF-κB response elements. Nature Communications, 2018, 9, 1337.	12.8	90
14	Defining a conformational ensemble that directs activation of PPAR $\hat{I}^3$ . Nature Communications, 2018, 9, 1794.	12.8	53
15	A structural mechanism for directing corepressor-selective inverse agonism of PPARÎ <sup>3</sup> . Nature Communications, 2018, 9, 4687.	12.8	38
16	Structural organization of a major neuronal G protein regulator, the RGS7-G $\hat{I}^2$ 5-R7BP complex. ELife, 2018, 7, .	6.0	18
17	REV-ERBα Regulates TH17 Cell Development and Autoimmunity. Cell Reports, 2018, 25, 3733-3749.e8.	6.4	78
18	Chemical Crosslinking Mass Spectrometry Reveals the Conformational Landscape of the Activation Helix of PPARÎ <sup>3</sup> ; a Model for Ligand-Dependent Antagonism. Structure, 2018, 26, 1431-1439.e6.	3.3	24

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19	Cooperative cobinding of synthetic and natural ligands to the nuclear receptor PPARÎ <sup>3</sup> . ELife, 2018, 7, .	6.0	53
20	Modification of the Orthosteric PPARÎ <sup>3</sup> Covalent Antagonist Scaffold Yields an Improved Dual-Site Allosteric Inhibitor. ACS Chemical Biology, 2017, 12, 969-978.	3.4	36
21	Systems Structural Biology Analysis of Ligand Effects on ERα Predicts Cellular Response to Environmental Estrogens and Anti-hormone Therapies. Cell Chemical Biology, 2017, 24, 35-45.	5.2	34
22	Tethering not required: the glucocorticoid receptor binds directly to activator protein-1 recognition motifs to repress inflammatory genes. Nucleic Acids Research, 2017, 45, 8596-8608.	14.5	69
23	Synergistic Regulation of Coregulator/Nuclear Receptor Interaction by Ligand and DNA. Structure, 2017, 25, 1506-1518.e4.	3.3	45
24	Identification of a Binding Site for Unsaturated Fatty Acids in the Orphan Nuclear Receptor Nurr1. ACS Chemical Biology, 2016, 11, 1795-1799.	3.4	59
25	Mechanistic insight into protein modification and sulfur mobilization activities of noncanonical E1 and associated ubiquitinâ€like proteins of Archaea. FEBS Journal, 2016, 283, 3567-3586.	4.7	21
26	Activity-Based Profiling Reveals a Regulatory Link between Oxidative Stress and Protein Arginine Phosphorylation. Cell Chemical Biology, 2016, 23, 967-977.	5.2	42
27	Probing the Complex Binding Modes of the PPARÎ <sup>3</sup> Partial Agonist 2-Chloro- <i>N</i> -(3-chloro-4-((5-chlorobenzo[ <i>d</i> )]thiazol-2-yl)thio)phenyl)-4-(trifluoromethyl)benzenesulfor (T2384) to Orthosteric and Allosteric Sites with NMR Spectroscopy. Journal of Medicinal Chemistry, 2016, 59, 10335-10341.	namide 6.4	24
28	Ebselen, a Small-Molecule Capsid Inhibitor of HIV-1 Replication. Antimicrobial Agents and Chemotherapy, 2016, 60, 2195-2208.	3.2	91
29	Distal substitutions drive divergent DNA specificity among paralogous transcription factors through subdivision of conformational space. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, 326-331.	7.1	28
30	Pharmacological repression of PPAR $\hat{I}^3$ promotes osteogenesis. Nature Communications, 2015, 6, 7443.	12.8	99
31	Anti-proliferative actions of a synthetic REV-ERBÎ $\pm$ /Î $^2$ agonist in breast cancer cells. Biochemical Pharmacology, 2015, 96, 315-322.	4.4	59
32	Structural mechanism for signal transduction in RXR nuclear receptor heterodimers. Nature Communications, 2015, 6, 8013.	12.8	101
33	Deconvolution of Complex 1D NMR Spectra Using Objective Model Selection. PLoS ONE, 2015, 10, e0134474.	2.5	15
34	Resveratrol modulates the inflammatory response via an estrogen receptor-signal integration network. ELife, 2014, 3, e02057.	6.0	113
35	Structure of REV-ERB $\hat{I}^2$ Ligand-binding Domain Bound to a Porphyrin Antagonist. Journal of Biological Chemistry, 2014, 289, 20054-20066.	3.4	22
36	REV-ERB and ROR nuclear receptors as drug targets. Nature Reviews Drug Discovery, 2014, 13, 197-216.	46.4	437

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37	Conserved sequence-specific lincRNA–steroid receptor interactions drive transcriptional repression and direct cell fate. Nature Communications, 2014, 5, 5395.	12.8	103
38	An alternate binding site for PPARγ ligands. Nature Communications, 2014, 5, 3571.	12.8	148
39	Conformational Allostery in Nuclear Receptor/Coregulator Transcriptional Complexes. Biophysical Journal, 2014, 106, 686a.	0.5	0
40	Ligand-binding dynamics rewire cellular signaling via estrogen receptor-α. Nature Chemical Biology, 2013, 9, 326-332.	8.0	53
41	Nuclear Receptors and Their Selective Pharmacologic Modulators. Pharmacological Reviews, 2013, 65, 710-778.	16.0	207
42	Small Molecule Modulation of Nuclear Receptor Conformational Dynamics: Implications for Function and Drug Discovery. Molecular Pharmacology, 2013, 83, 1-8.	2.3	100
43	Small molecule tertiary amines as agonists of the nuclear hormone receptor Rev-erbl±. Bioorganic and Medicinal Chemistry Letters, 2012, 22, 4413-4417.	2.2	16
44	Regulation of circadian behaviour and metabolism by synthetic REV-ERB agonists. Nature, 2012, 485, 62-68.	27.8	638
45	1H, 13C and 15N chemical shift assignments for the human Pitx2 homeodomain in complex with a 22-base hairpin DNA. Biomolecular NMR Assignments, 2012, 6, 79-81.	0.8	0
46	Synthesis and SAR of tetrahydroisoquinolines as Rev-erbα agonists. Bioorganic and Medicinal Chemistry Letters, 2012, 22, 3739-3742.	2.2	22
47	Ligand and Receptor Dynamics Contribute to the Mechanism of Graded PPARÎ <sup>3</sup> Agonism. Structure, 2012, 20, 139-150.	3.3	133
48	Regulation of p53 Stability and Apoptosis by a ROR Agonist. PLoS ONE, 2012, 7, e34921.	2.5	54
49	Identification of SR8278, a Synthetic Antagonist of the Nuclear Heme Receptor REV-ERB. ACS Chemical Biology, 2011, 6, 131-134.	3.4	152
50	Identification of SR3335 (ML-176): A Synthetic RORα Selective Inverse Agonist. ACS Chemical Biology, 2011, 6, 218-222.	3.4	114
51	DNA binding alters coactivator interaction surfaces of the intact VDR–RXR complex. Nature Structural and Molecular Biology, 2011, 18, 556-563.	8.2	185
52	Observing selected domains in multi-domain proteins via sortase-mediated ligation and NMR spectroscopy. Journal of Biomolecular NMR, 2011, 49, 3-7.	2.8	40
53	1H, 13C and 15N chemical shift assignments for the human Pitx2 homeodomain and a R24H homeodomain mutant. Biomolecular NMR Assignments, 2011, 5, 105-107.	0.8	4
54	The REV-ERBs and RORs: molecular links between circadian rhythms and lipid homeostasis. Future Medicinal Chemistry, 2011, 3, 623-638.	2.3	131

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55	Characterization of the Core Mammalian Clock Component, NPAS2, as a REV-ERBα/RORα Target Gene. Journal of Biological Chemistry, 2010, 285, 35386-35392.	3.4	117
56	Structural and Motional Contributions of the Bacillus subtilis ClpC N-Domain to Adaptor Protein Interactions. Journal of Molecular Biology, 2009, 387, 639-652.	4.2	18
57	Insights into the Nature of DNA Binding of AbrB-like Transcription Factors. Structure, 2008, 16, 1702-1713.	3.3	30
58	Implications of the binding of tamoxifen to the coactivator recognition site of the estrogen receptor. Endocrine-Related Cancer, 2008, 15, 851-870.	3.1	49
59	Classification of Response Regulators Based on Their Surface Properties. Methods in Enzymology, 2007, 422, 141-169.	1.0	2
60	NMR assignment of the N-terminal repeat domain of Bacillus subtilis ClpC. Biomolecular NMR Assignments, 2007, $1,163-165$ .	0.8	2
61	Structure, binding interface and hydrophobic transitions of Ca2+-loaded calbindin-D28K. Nature Structural and Molecular Biology, 2006, 13, 641-647.	8.2	75
62	Structural Analysis of Divalent Metals Binding to the Bacillus subtilis Response Regulator Spo0F: The Possibility for In Vitro Metalloregulation in the Initiation of Sporulation. BioMetals, 2005, 18, 449-466.	4.1	19
63	Solution Structure and Dynamics of LuxU from Vibrio harveyi, a Phosphotransferase Protein Involved in Bacterial Quorum Sensing. Journal of Molecular Biology, 2005, 347, 297-307.	4.2	20
64	Corrigendum to: Sub-classification of response regulators using the surface characteristics of their receiver domains (FEBS 27785). FEBS Letters, 2004, 560, 227-228.	2.8	1
65	Sub-classification of response regulators using the surface characteristics of their receiver domains. FEBS Letters, 2003, 554, 231-236.	2.8	7
66	Alternative Splicing of a $\hat{l}^24$ Subunit Proline-Rich Motif Regulates Voltage-Dependent Gating and Toxin Block of Cav2.1 Ca2+Channels. Journal of Neuroscience, 2002, 22, 9331-9339.	3.6	25