Dino Moras

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/970589/publications.pdf

Version: 2024-02-01

22 papers 4,722 citations

471509 17 h-index 677142 22 g-index

23 all docs 23 docs citations

times ranked

23

2441 citing authors

#	Article	IF	CITATIONS
1	Crystal structure of the ligand-binding domain of the human nuclear receptor RXR-α. Nature, 1995, 375, 377-382.	27.8	1,155
2	Crystal structure of the RAR- \hat{I}^3 ligand-binding domain bound to all-trans retinoic acid. Nature, 1995, 378, 681-689.	27.8	1,115
3	A canonical structure for the ligand-binding domain of nuclear receptors. Nature Structural Biology, 1996, 3, 87-94.	9.7	859
4	Crystal Structure of a Heterodimeric Complex of RAR and RXR Ligand-Binding Domains. Molecular Cell, 2000, 5, 289-298.	9.7	385
5	Structural adaptability in the ligand-binding pocket of the ecdysone hormone receptor. Nature, 2003, 426, 91-96.	27.8	239
6	Crystal Structure of the Ligand-binding Domain of the Ultraspiracle Protein USP, the Ortholog of Retinoid X Receptors in Insects. Journal of Biological Chemistry, 2001, 276, 7465-7474.	3.4	157
7	Molecular Recognition of Agonist Ligands by RXRs. Molecular Endocrinology, 2002, 16, 987-997.	3.7	154
8	Structural and functional characterization of a novel type of ligand-independent RXR-USP receptor. EMBO Journal, 2007, 26, 3770-3782.	7.8	120
9	Adaptability of the Vitamin D nuclear receptor to the synthetic ligand Gemini: Remodelling the LBP with one side chain rotation. Journal of Steroid Biochemistry and Molecular Biology, 2007, 103, 235-242.	2.5	85
10	Signature of the oligomeric behaviour of nuclear receptors at the sequence and structural level. EMBO Reports, 2004, 5, 423-429.	4.5	80
11	Allosteric Controls of Nuclear Receptor Function in the Regulation of Transcription. Journal of Molecular Biology, 2013, 425, 2317-2329.	4.2	55
12	Structural analysis of nuclear receptors: From isolated domains to integral proteins. Molecular and Cellular Endocrinology, 2012, 348, 466-473.	3.2	54
13	Molecular Recognition of Agonist Ligands by RXRs. Molecular Endocrinology, 2002, 16, 987-997.	3.7	48
14	Structural and Evolutionary Innovation of the Heterodimerization Interface between USP and the Ecdysone Receptor ECR in Insects. Molecular Biology and Evolution, 2009, 26, 753-768.	8.9	45
15	A Vitamin D Receptor Selectively Activated by Gemini Analogs Reveals Ligand Dependent and Independent Effects. Cell Reports, 2015, 10, 516-526.	6.4	45
16	The Asymmetric Binding of PGC-1α to the ERRα and ERRγ Nuclear Receptor Homodimers Involves a Similar Recognition Mechanism. PLoS ONE, 2013, 8, e67810.	2.5	34
17	The palindromic DNA-bound USP/EcR nuclear receptor adopts an asymmetric organization with allosteric domain positioning. Nature Communications, 2014, 5, 4139.	12.8	33
18	Structural Insights into the Polyphyletic Origins of Glycyl tRNA Synthetases. Journal of Biological Chemistry, 2016, 291, 14430-14446.	3.4	16

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#	Article	IF	CITATION
19	A revisited version of the apo structure of the ligand-binding domain of the human nuclear receptor retinoic X receptor \hat{l}_{\pm} . Acta Crystallographica Section F, Structural Biology Communications, 2019, 75, 98-104.	0.8	14
20	Structure–function relationships in nuclear receptors: the facts. Trends in Biochemical Sciences, 2015, 40, 287-290.	7.5	12
21	The dual role of CHAPS in the crystallization of stromelysin-3 catalytic domain. Acta Crystallographica Section D: Biological Crystallography, 2003, 59, 603-606.	2.5	9
22	A structural signature motif enlightens the origin and diversification of nuclear receptors. PLoS Genetics, 2021, 17, e1009492.	3.5	8