

Albane le Maire

List of Publications by Year in descending order

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24
papers

1,675
citations

516215

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642321

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docs citations

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times ranked

2902
citing authors

#	ARTICLE	IF	CITATIONS
1	Retinoids. , 2021, , 1358-1367.		0
2	PPAR β S273 Phosphorylation Modifies the Dynamics of Coregulator Proteins Recruitment. <i>Frontiers in Endocrinology</i> , 2020, 11, 561256.	1.5	7
3	Protein-protein interactions in the regulation of RAR α -RXR heterodimers transcriptional activity. <i>Methods in Enzymology</i> , 2020, 637, 175-207.	0.4	5
4	Two Novel Cases of Resistance to Thyroid Hormone Due to <i>THRA</i> Mutation. <i>Thyroid</i> , 2020, 30, 1217-1221.	2.4	16
5	Interplay of Protein Disorder in Retinoic Acid Receptor Heterodimer and Its Corepressor Regulates Gene Expression. <i>Structure</i> , 2019, 27, 1270-1285.e6.	1.6	50
6	Regulation of RXR-RAR Heterodimers by RXR- and RAR-Specific Ligands and Their Combinations. <i>Cells</i> , 2019, 8, 1392.	1.8	55
7	Pathological Interactions Between Mutant Thyroid Hormone Receptors and Corepressors and Their Modulation by a Thyroid Hormone Analogue with Therapeutic Potential. <i>Thyroid</i> , 2018, 28, 1708-1722.	2.4	9
8	Screening for PPAR Non-Agonist Ligands Followed by Characterization of a Hit, AM-879, with Additional No-Adipogenic and cdk5-Mediated Phosphorylation Inhibition Properties. <i>Frontiers in Endocrinology</i> , 2018, 9, 11.	1.5	21
9	The Human Mixed Lineage Leukemia 5 (MLL5), a Sequentially and Structurally Divergent SET Domain-Containing Protein with No Intrinsic Catalytic Activity. <i>PLoS ONE</i> , 2016, 11, e0165139.	1.1	31
10	Molecular mechanisms of transcriptional control by Rev β : An energetic foundation for reconciling structure and binding with biological function. <i>Protein Science</i> , 2015, 24, 1129-1146.	3.1	11
11	Nuclear Receptor Profiling of Bisphenol-A and Its Halogenated Analogues. <i>Vitamins and Hormones</i> , 2014, 94, 229-251.	0.7	59
12	Retinoic Acid Receptors: Structural Basis for Coregulator Interaction and Exchange. <i>Sub-Cellular Biochemistry</i> , 2014, 70, 37-54.	1.0	27
13	Retinoid Receptors and Therapeutic Applications of RAR/RXR Modulators. <i>Current Topics in Medicinal Chemistry</i> , 2012, 12, 505-527.	1.0	86
14	Structural and mechanistic insights into bisphenols action provide guidelines for risk assessment and discovery of bisphenol A substitutes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 14930-14935.	3.3	313
15	In-plate protein crystallization, <i>in situ</i> ligand soaking and X-ray diffraction. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2011, 67, 747-755.	2.5	70
16	Characterization of Novel Ligands of ER α , ER β , and PPAR β : The Case of Halogenated Bisphenol A and Their Conjugated Metabolites. <i>Toxicological Sciences</i> , 2011, 122, 372-382.	1.4	119
17	Peroxisome Proliferator-Activated Receptor β Is a Target for Halogenated Analogs of Bisphenol A. <i>Environmental Health Perspectives</i> , 2011, 119, 1227-1232.	2.8	257
18	A structural view of nuclear hormone receptor: endocrine disruptor interactions. <i>Cellular and Molecular Life Sciences</i> , 2010, 67, 1219-1237.	2.4	105

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19	A unique secondary-structure switch controls constitutive gene repression by retinoic acid receptor. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 801-807.	3.6	142
20	Activation of RXR-PPAR heterodimers by organotin environmental endocrine disruptors. <i>EMBO Reports</i> , 2009, 10, 367-373.	2.0	235
21	Solution structure of the region 51-160 of human KIN17 reveals an atypical winged helix domain. <i>Protein Science</i> , 2007, 16, 2750-2755.	3.1	20
22	A Tandem of SH3-like Domains Participates in RNA Binding in KIN17, a Human Protein Activated in Response to Genotoxics. <i>Journal of Molecular Biology</i> , 2006, 364, 764-776.	2.0	20
23	Crystallization and halide phasing of the C-terminal domain of human KIN17. <i>Acta Crystallographica Section F: Structural Biology Communications</i> , 2006, 62, 245-248.	0.7	7
24	Solution NMR structure of the SH3 domain of human nephrocystin and analysis of a mutation-causing juvenile nephronophthisis. <i>Proteins: Structure, Function and Bioinformatics</i> , 2005, 59, 347-355.	1.5	10