## Colin W Levy

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7048611/publications.pdf

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50	1,929	22	42
papers	citations	h-index	g-index
50	50	50	2556
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Reductive dehalogenase structure suggests a mechanism for B12-dependent dehalogenation. Nature, 2015, 517, 513-516.	27.8	260
2	Better than Nature: Nicotinamide Biomimetics That Outperform Natural Coenzymes. Journal of the American Chemical Society, 2016, 138, 1033-1039.	13.7	164
3	The road to fully programmable protein catalysis. Nature, 2022, 606, 49-58.	27.8	126
4	Extending the biocatalytic scope of regiocomplementary flavin-dependent halogenase enzymes. Chemical Science, 2015, 6, 3454-3460.	7.4	89
5	Structural basis for enzymatic photocatalysis in chlorophyll biosynthesis. Nature, 2019, 574, 722-725.	27.8	88
6	Design and evolution of an enzyme with a non-canonical organocatalytic mechanism. Nature, 2019, 570, 219-223.	27.8	86
7	Updated structure of Drosophila cryptochrome. Nature, 2013, 495, E3-E4.	27.8	83
8	Catalytic Determinants of Alkene Production by the Cytochrome P450 Peroxygenase OleTJE. Journal of Biological Chemistry, 2017, 292, 5128-5143.	3.4	73
9	A Structureâ€Guided Switch in the Regioselectivity of a Tryptophan Halogenase. ChemBioChem, 2016, 17, 821-824.	2.6	71
10	Biocatalytic Asymmetric Alkene Reduction: Crystal Structure and Characterization of a Double Bond Reductase from <i>Nicotiana tabacum</i> . ACS Catalysis, 2013, 3, 370-379.	11.2	59
11	Effects of Activeâ€6ite Modification and Quaternary Structure on the Regioselectivity of Catecholâ€∢i>Oà6€Methyltransferase. Angewandte Chemie - International Edition, 2016, 55, 2683-2687.	13.8	58
12	The major secreted protein of the whipworm parasite tethers to matrix and inhibits interleukin-13 function. Nature Communications, 2019, 10, 2344.	12.8	48
13	Fragment-Based Approaches to the Development of <i>Mycobacterium tuberculosis</i> CYP121 Inhibitors. Journal of Medicinal Chemistry, 2016, 59, 3272-3302.	6.4	47
14	Discovery, characterization and engineering of ligases for amide synthesis. Nature, 2021, 593, 391-398.	27.8	40
15	Structure and Biocatalytic Scope of Coclaurine <i>N</i> â€Methyltransferase. Angewandte Chemie - International Edition, 2018, 57, 10600-10604.	13.8	37
16	A brain-permeable inhibitor of the neurodegenerative disease target kynurenine 3-monooxygenase prevents accumulation of neurotoxic metabolites. Communications Biology, 2019, 2, 271.	4.4	36
17	Engineering an efficient and enantioselective enzyme for the Morita–Baylis–Hillman reaction. Nature Chemistry, 2022, 14, 313-320.	13.6	34
18	Carboplatin binding to histidine. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 1135-1142.	0.8	33

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19	Discovery and Optimization of Allosteric Inhibitors of Mutant Isocitrate Dehydrogenase 1 (R132H IDH1) Displaying Activity in Human Acute Myeloid Leukemia Cells. Journal of Medicinal Chemistry, 2016, 59, 11120-11137.	6.4	31
20	Structures of <i>Arabidopsis thaliana</i> oxygen-sensing plant cysteine oxidases 4 and 5 enable targeted manipulation of their activity. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 23140-23147.	7.1	31
21	Novel Aryl Substituted Pyrazoles as Small Molecule Inhibitors of Cytochrome P450 CYP121A1: Synthesis and Antimycobacterial Evaluation. Journal of Medicinal Chemistry, 2017, 60, 10257-10267.	6.4	26
22	Effects of Activeâ€Site Modification and Quaternary Structure on the Regioselectivity of Catecholâ€ <i>O</i> ÀeMethyltransferase. Angewandte Chemie, 2016, 128, 2733-2737.	2.0	25
23	Rewiring the "Push-Pull―Catalytic Machinery of a Heme Enzyme Using an Expanded Genetic Code. ACS Catalysis, 2020, 10, 2735-2746.	11.2	25
24	The structure of the folded domain from the signature multifunctional protein ICP27 from herpes simplex virus-1 reveals an intertwined dimer. Scientific Reports, 2015, 5, 11234.	3.3	23
25	The herpes viral transcription factor ICP4 forms a novel DNA recognition complex. Nucleic Acids Research, 2017, 45, 8064-8078.	14.5	23
26	Structure and Properties of a Natural Competence-Associated Pilin Suggest a Unique Pilus Tip-Associated DNA Receptor. MBio, 2019, 10, .	4.1	23
27	Structural Basis for Selective Interaction between the ESCRT Regulator HD-PTP and UBAP1. Structure, 2016, 24, 2115-2126.	3.3	22
28	Equatorial Active Site Compaction and Electrostatic Reorganization in Catechol-O-methyltransferase. ACS Catalysis, 2019, 9, 4394-4401.	11.2	21
29	Structure and Mechanism of a Viral Collagen Prolyl Hydroxylase. Biochemistry, 2015, 54, 6093-6105.	2.5	19
30	Design and Synthesis of Imidazole and Triazole Pyrazoles as <i>Mycobacterium Tuberculosis</i> CYP121A1 Inhibitors. ChemistryOpen, 2019, 8, 995-1011.	1.9	19
31	Structure–Activity Relationships of <i>cyclo</i> ((scp>l-Tyrosyl- <scp>l</scp> -tyrosine) Derivatives Binding to <i>Mycobacterium tuberculosis</i> CYP121: Iodinated Analogues Promote Shift to High-Spin Adduct. Journal of Medicinal Chemistry, 2019, 62, 9792-9805.	6.4	19
32	Structural and catalytic properties of the peroxygenase P450 enzyme CYP152K6 from Bacillus methanolicus. Journal of Inorganic Biochemistry, 2018, 188, 18-28.	3.5	18
33	Novel insights into P450 BM3 interactions with FDA-approved antifungal azole drugs. Scientific Reports, 2019, 9, 1577.	3.3	17
34	Molecular Mechanism of SR Protein Kinase 1 Inhibition by the Herpes Virus Protein ICP27. MBio, 2019, 10,	4.1	17
35	UbiD domain dynamics underpins aromatic decarboxylation. Nature Communications, 2021, 12, 5065.	12.8	14
36	Structural Characterization and Ligand/Inhibitor Identification Provide Functional Insights into the Mycobacterium tuberculosis Cytochrome P450 CYP126A1. Journal of Biological Chemistry, 2017, 292, 1310-1329.	3.4	13

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37	Design, synthesis and evaluation against Mycobacterium tuberculosis of azole piperazine derivatives as dicyclotyrosine (cYY) mimics. Bioorganic and Medicinal Chemistry, 2018, 26, 161-176.	3.0	13
38	The crystal structure of P450-TT heme-domain provides the first structural insights into the versatile class VII P450s. Biochemical and Biophysical Research Communications, 2018, 501, 846-850.	2.1	13
39	Nonequivalence of Second Sphere "Noncatalytic―Residues in Pentaerythritol Tetranitrate Reductase in Relation to Local Dynamics Linked to H-Transfer in Reactions with NADH and NADPH Coenzymes. ACS Catalysis, 2018, 8, 11589-11599.	11.2	12
40	Structural basis of terephthalate recognition by solute binding protein TphC. Nature Communications, 2021, 12, 6244.	12.8	12
41	Response from Tanley <i>et al.</i> to <i>Crystallography and chemistry should always go together: a cautionary tale of protein complexes with cisplatin and carboplatin</i> . Acta Crystallographica Section D: Biological Crystallography, 2015, 71, 1982-1983.	2.5	11
42	Structural and biochemical characterization of the prenylated flavin mononucleotide-dependent indole-3-carboxylic acid decarboxylase. Journal of Biological Chemistry, 2022, 298, 101771.	3.4	10
43	The Dual PDZ Domain from Postsynaptic Density Protein 95 Forms a Scaffold with Peptide Ligand. Biophysical Journal, 2020, 119, 667-689.	0.5	9
44	A Noncanonical Tryptophan Analogue Reveals an Active Site Hydrogen Bond Controlling Ferryl Reactivity in a Heme Peroxidase. Jacs Au, 2021, 1, 913-918.	7.9	8
45	Structural characterization of CYP144A1 – a cytochrome P450 enzyme expressed from alternative transcripts in Mycobacterium tuberculosis. Scientific Reports, 2016, 6, 26628.	3.3	7
46	Structure and Biocatalytic Scope of Coclaurine N â€Methyltransferase. Angewandte Chemie, 2018, 130, 10760-10764.	2.0	6
47	Structural identification of conserved RNA binding sites in herpesvirus ORF57 homologs: implications for PAN RNA recognition. Nucleic Acids Research, 2019, 47, 1987-2001.	14.5	4
48	A new strategy for hit generation: Novel in cellulo active inhibitors of CYP121A1 from Mycobacterium tuberculosis via a combined X-ray crystallographic and phenotypic screening approach (XP screen). European Journal of Medicinal Chemistry, 2022, 230, 114105.	5.5	4
49	Studies of the oligomerisation mechanism of a cystatin-based engineered protein scaffold. Scientific Reports, 2019, 9, 9067.	3.3	2
50	Crystallization of PTP Domains. Methods in Molecular Biology, 2016, 1447, 155-180.	0.9	0