## 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  | 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         |
| 2  | 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        |
| 3  | Engineering an efficient and enantioselective enzyme for the Morita–Baylis–Hillman reaction. Nature<br>Chemistry, 2022, 14, 313-320.   | 13.6 | 34        |
| 4  | The road to fully programmable protein catalysis. Nature, 2022, 606, 49-58.  | 27.8 | 126       |
| 5  | Discovery, characterization and engineering of ligases for amide synthesis. Nature, 2021, 593, 391-398.  | 27.8 | 40        |
| 6  | 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         |
| 7  | UbiD domain dynamics underpins aromatic decarboxylation. Nature Communications, 2021, 12, 5065.  | 12.8 | 14        |
| 8  | Structural basis of terephthalate recognition by solute binding protein TphC. Nature Communications, 2021, 12, 6244.   | 12.8 | 12        |
| 9  | 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        |
| 10 | The Dual PDZ Domain from Postsynaptic Density Protein 95 Forms a Scaffold with Peptide Ligand. Biophysical Journal, 2020, 119, 667-689.  | 0.5  | 9         |
| 11 | Rewiring the "Push-Pull―Catalytic Machinery of a Heme Enzyme Using an Expanded Genetic Code. ACS<br>Catalysis, 2020, 10, 2735-2746.  | 11.2 | 25        |
| 12 | Design and Synthesis of Imidazole and Triazole Pyrazoles as <i>Mycobacterium Tuberculosis</i> CYP121A1 Inhibitors. ChemistryOpen, 2019, 8, 995-1011.   | 1.9  | 19        |
| 13 | 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        |
| 14 | Structure–Activity Relationships of <i>cyclo</i> ( <scp> </scp> -Tyrosyl- <scp> </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        |
| 15 | Studies of the oligomerisation mechanism of a cystatin-based engineered protein scaffold. Scientific Reports, 2019, 9, 9067.   | 3.3  | 2         |
| 16 | The major secreted protein of the whipworm parasite tethers to matrix and inhibits interleukin-13 function. Nature Communications, 2019, 10, 2344.   | 12.8 | 48        |
| 17 | Design and evolution of an enzyme with a non-canonical organocatalytic mechanism. Nature, 2019, 570, 219-223.  | 27.8 | 86        |
| 18 | Structure and Properties of a Natural Competence-Associated Pilin Suggest a Unique Pilus Tip-Associated DNA Receptor. MBio, 2019, 10, .  | 4.1  | 23        |

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|----|--|------|-----------|
| 19 | Equatorial Active Site Compaction and Electrostatic Reorganization in Catechol-O-methyltransferase. ACS Catalysis, 2019, 9, 4394-4401.   | 11.2 | 21        |
| 20 | Novel insights into P450 BM3 interactions with FDA-approved antifungal azole drugs. Scientific Reports, 2019, 9, 1577.   | 3.3  | 17        |
| 21 | Molecular Mechanism of SR Protein Kinase 1 Inhibition by the Herpes Virus Protein ICP27. MBio, 2019, 10,   | 4.1  | 17        |
| 22 | Structural basis for enzymatic photocatalysis in chlorophyll biosynthesis. Nature, 2019, 574, 722-725.   | 27.8 | 88        |
| 23 | 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         |
| 24 | 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        |
| 25 | Nonequivalence of Second Sphere "Noncatalytic―Residues in Pentaerythritol Tetranitrate Reductase<br>in Relation to Local Dynamics Linked to H-Transfer in Reactions with NADH and NADPH Coenzymes. ACS<br>Catalysis, 2018, 8, 11589-11599. | 11.2 | 12        |
| 26 | Structure and Biocatalytic Scope of Coclaurine <i>N</i> â€Methyltransferase. Angewandte Chemie - International Edition, 2018, 57, 10600-10604.   | 13.8 | 37        |
| 27 | 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        |
| 28 | Structural and catalytic properties of the peroxygenase P450 enzyme CYP152K6 from Bacillus methanolicus. Journal of Inorganic Biochemistry, 2018, 188, 18-28.  | 3.5  | 18        |
| 29 | Structure and Biocatalytic Scope of Coclaurine N â€Methyltransferase. Angewandte Chemie, 2018, 130, 10760-10764.   | 2.0  | 6         |
| 30 | Catalytic Determinants of Alkene Production by the Cytochrome P450 Peroxygenase OleTJE. Journal of Biological Chemistry, 2017, 292, 5128-5143.   | 3.4  | 73        |
| 31 | The herpes viral transcription factor ICP4 forms a novel DNA recognition complex. Nucleic Acids Research, 2017, 45, 8064-8078.   | 14.5 | 23        |
| 32 | 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        |
| 33 | 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        |
| 34 | Structural Basis for Selective Interaction between the ESCRT Regulator HD-PTP and UBAP1. Structure, 2016, 24, 2115-2126.   | 3.3  | 22        |
| 35 | Effects of Activeâ€Site Modification and Quaternary Structure on the Regioselectivity of Catecholâ€∢i>O∢/i>â€Methyltransferase. Angewandte Chemie - International Edition, 2016, 55, 2683-2687.  | 13.8 | 58        |
| 36 | A Structureâ€Guided Switch in the Regioselectivity of a Tryptophan Halogenase. ChemBioChem, 2016, 17, 821-824.   | 2.6  | 71        |

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|----|--|------|----------|
| 37 | Structural characterization of CYP144A1 – a cytochrome P450 enzyme expressed from alternative transcripts in Mycobacterium tuberculosis. Scientific Reports, 2016, 6, 26628.   | 3.3  | 7        |
| 38 | 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       |
| 39 | Crystallization of PTP Domains. Methods in Molecular Biology, 2016, 1447, 155-180.   | 0.9  | 0        |
| 40 | Effects of Activeâ€Site Modification and Quaternary Structure on the Regioselectivity of Catecholâ€∢i>Oàâ€Methyltransferase. Angewandte Chemie, 2016, 128, 2733-2737.  | 2.0  | 25       |
| 41 | Fragment-Based Approaches to the Development of <i>Mycobacterium tuberculosis</i> CYP121 Inhibitors. Journal of Medicinal Chemistry, 2016, 59, 3272-3302.  | 6.4  | 47       |
| 42 | Better than Nature: Nicotinamide Biomimetics That Outperform Natural Coenzymes. Journal of the American Chemical Society, 2016, 138, 1033-1039.  | 13.7 | 164      |
| 43 | 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       |
| 44 | Extending the biocatalytic scope of regiocomplementary flavin-dependent halogenase enzymes. Chemical Science, 2015, 6, 3454-3460.  | 7.4  | 89       |
| 45 | Structure and Mechanism of a Viral Collagen Prolyl Hydroxylase. Biochemistry, 2015, 54, 6093-6105.   | 2.5  | 19       |
| 46 | Reductive dehalogenase structure suggests a mechanism for B12-dependent dehalogenation. Nature, 2015, 517, 513-516.  | 27.8 | 260      |
| 47 | 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       |
| 48 | Carboplatin binding to histidine. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 1135-1142.  | 0.8  | 33       |
| 49 | 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       |
| 50 | Undated structure of Drosophila cryptochrome Nature 2013 495 F3-F4   | 27.8 | 83       |