Axel Imhof

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8614363/publications.pdf

Version: 2024-02-01

200 papers 17,264 citations

19657 61 h-index 17105 122 g-index

221 all docs

221 docs citations

times ranked

221

21659 citing authors

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | ImShot: An Open-Source Software for Probabilistic Identification of Proteins In Situ and Visualization of Proteomics Data. Molecular and Cellular Proteomics, 2022, 21, 100242. | 3.8 | 3 |
| 2 | Dietary intervention improves health metrics and life expectancy of the genetically obese Titan mouse. Communications Biology, 2022, 5, 408. | 4.4 | 4 |
| 3 | Endotoxinemia Accelerates Atherosclerosis Through Electrostatic Charge–Mediated Monocyte Adhesion. Circulation, 2021, 143, 254-266. | 1.6 | 266 |
| 4 | Reduced peroxisomal import triggers peroxisomal retrograde signaling. Cell Reports, 2021, 34, 108653. | 6.4 | 9 |
| 5 | <i>Helicobacter hepaticus</i> is required for immune targeting of bacterial heat shock protein 60 and fatal colitis in mice. Gut Microbes, 2021, 13, 1-20. | 9.8 | 8 |
| 6 | Metabolic Analysis of Vitreous/Lens and Retina in Wild Type and Retinal Degeneration Mice. International Journal of Molecular Sciences, 2021, 22, 2345. | 4.1 | 6 |
| 7 | A novel proteomics approach to epigenetic profiling of circulating nucleosomes. Scientific Reports, 2021, 11, 7256. | 3.3 | 21 |
| 8 | Proteome dynamics at broken replication forks reveal a distinct ATM-directed repair response suppressing DNA double-strand break ubiquitination. Molecular Cell, 2021, 81, 1084-1099.e6. | 9.7 | 57 |
| 9 | Exploring the Ion Channel TRPV2 and Testicular Macrophages in Mouse Testis. International Journal of Molecular Sciences, 2021, 22, 4727. | 4.1 | 5 |
| 10 | Systematic functional analysis of SARS-CoV-2 proteins uncovers viral innate immune antagonists and remaining vulnerabilities. Cell Reports, 2021, 35, 109126. | 6.4 | 176 |
| 11 | Investigation and Highly Accurate Prediction of Missed Tryptic Cleavages by Deep Learning. Journal of Proteome Research, 2021, 20, 3749-3757. | 3.7 | 9 |
| 12 | Pumilio2 and Staufen2 selectively balance the synaptic proteome. Cell Reports, 2021, 35, 109279. | 6.4 | 14 |
| 13 | Phosphorylation of the HP1 \hat{l}^2 hinge region sequesters KAP1 in heterochromatin and promotes the exit from na \hat{A} ve pluripotency. Nucleic Acids Research, 2021, 49, 7406-7423. | 14.5 | 9 |
| 14 | GSNOR Contributes to Demethylation and Expression of Transposable Elements and Stress-Responsive Genes. Antioxidants, 2021, 10, 1128. | 5.1 | 10 |
| 15 | The Integrity of the HMR complex is necessary for centromeric binding and reproductive isolation in Drosophila. PLoS Genetics, 2021, 17, e1009744. | 3.5 | 35 |
| 16 | A systemic cell cycle block impacts stage-specific histone modification profiles during Xenopus embryogenesis. PLoS Biology, 2021, 19, e3001377. | 5.6 | 2 |
| 17 | MALDI-IMS combined with shotgun proteomics identify and localize new factors in male infertility. Life Science Alliance, 2021, 4, e202000672. | 2.8 | 7 |
| 18 | Discovery of Native Protein Complexes by Liquid Chromatography Followed byÂQuantitative Mass Spectrometry. Advances in Experimental Medicine and Biology, 2021, 1336, 105-128. | 1.6 | 0 |

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| 19 | Morc3 silences endogenous retroviruses by enabling Daxx-mediated histone H3.3 incorporation. Nature Communications, 2021, 12, 5996. | 12.8 | 34 |
| 20 | Determining histone H4 acetylation patterns in human peripheral blood mononuclear cells using mass spectrometry. Clinical Mass Spectrometry, 2020, 15, 54-60. | 1.9 | 3 |
| 21 | Molecular Wiring of a Mitochondrial Translational Feedback Loop. Molecular Cell, 2020, 77, 887-900.e5. | 9.7 | 22 |
| 22 | BMAL1 Associates with NOP58 in the Nucleolus and Contributes to Pre-rRNA Processing. IScience, 2020, 23, 101151. | 4.1 | 13 |
| 23 | S-adenosyl- <scp> </scp> -homocysteine hydrolase links methionine metabolism to the circadian clock and chromatin remodeling. Science Advances, 2020, 6, . | 10.3 | 49 |
| 24 | H4K20 Methylation Is Differently Regulated by Dilution and Demethylation in Proliferating and Cell-Cycle-Arrested Xenopus Embryos. Cell Systems, 2020, 11, 653-662.e8. | 6.2 | 6 |
| 25 | Histone Modifications in Stem Cell Development and Their Clinical Implications. Stem Cell Reports, 2020, 15, 1196-1205. | 4.8 | 17 |
| 26 | Physical Activity Dynamically Regulates the Hippocampal Proteome along the Dorso-Ventral Axis. International Journal of Molecular Sciences, 2020, 21, 3501. | 4.1 | 4 |
| 27 | Spt6 is a maintenance factor for centromeric CENP-A. Nature Communications, 2020, 11, 2919. | 12.8 | 30 |
| 28 | Harmonization of quality metrics and power calculation in multi-omic studies. Nature Communications, 2020, 11, 3092. | 12.8 | 43 |
| 29 | A multi-layered structure of the interphase chromocenter revealed by proximity-based biotinylation. Nucleic Acids Research, 2020, 48, 4161-4178. | 14.5 | 11 |
| 30 | Chromosome organization by a conserved condensin-ParB system in the actinobacterium Corynebacterium glutamicum. Nature Communications, 2020, 11, 1485. | 12.8 | 64 |
| 31 | Structure and Function of an Elongation Factor P Subfamily in Actinobacteria. Cell Reports, 2020, 30, 4332-4342.e5. | 6.4 | 11 |
| 32 | Msp1 cooperates with the proteasome for extraction of arrested mitochondrial import intermediates. Molecular Biology of the Cell, 2020, 31, 753-767. | 2.1 | 32 |
| 33 | Mechanisms governing the pioneering and redistribution capabilities of the non-classical pioneer PU.1. Nature Communications, 2020, 11 , 402. | 12.8 | 76 |
| 34 | New Approaches for Absolute Quantification of Stableâ€Isotopeâ€Labeled Peptide Standards for Targeted Proteomics Based on a UV Active Tag. Proteomics, 2020, 20, e2000007. | 2.2 | 7 |
| 35 | Domain Model Explains Propagation Dynamics and Stability of Histone H3K27 and H3K36 Methylation Landscapes. Cell Reports, 2020, 30, 1223-1234.e8. | 6.4 | 54 |
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| 37 | Trnp1 organizes diverse nuclear membraneâ€less compartments in neural stem cells. EMBO Journal, 2020, 39, e103373. | 7.8 | 16 |
| 38 | Mapping protein networks in yeast mitochondria using proximity-dependent biotin identification coupled to proteomics. STAR Protocols, 2020, 1, 100219. | 1.2 | 1 |
| 39 | The Impact of One Carbon Metabolism on Histone Methylation. Frontiers in Genetics, 2019, 10, 764. | 2.3 | 72 |
| 40 | Quantification of Proteins and Histone Marks in Drosophila Embryos Reveals Stoichiometric Relationships Impacting Chromatin Regulation. Developmental Cell, 2019, 51, 632-644.e6. | 7.0 | 50 |
| 41 | STATegra, a comprehensive multi-omics dataset of B-cell differentiation in mouse. Scientific Data, 2019, 6, 256. | 5.3 | 26 |
| 42 | Distinct CoREST complexes act in a cell-type-specific manner. Nucleic Acids Research, 2019, 47, 11649-11666. | 14.5 | 10 |
| 43 | A Drosophila cell-free system that senses DNA breaks and triggers phosphorylation signalling. Nucleic Acids Research, 2019, 47, 7444-7459. | 14.5 | 4 |
| 44 | Altered Localization of Hybrid Incompatibility Proteins in Drosophila. Molecular Biology and Evolution, 2019, 36, 1783-1792. | 8.9 | 9 |
| 45 | SETDB1-dependent heterochromatin stimulates alternative lengthening of telomeres. Science Advances, 2019, 5, eaav3673. | 10.3 | 70 |
| 46 | KMT9 monomethylates histone H4 lysine 12 and controls proliferation of prostate cancer cells. Nature Structural and Molecular Biology, 2019, 26, 361-371. | 8.2 | 57 |
| 47 | Nucleoside analogue activators of cyclic AMP-independent protein kinase A of Trypanosoma. Nature Communications, 2019, 10, 1421. | 12.8 | 33 |
| 48 | Multi-Reference Spectral Library Yields Almost Complete Coverage of Heterogeneous LC-MS/MS Data Sets. Journal of Proteome Research, 2019, 18, 1553-1566. | 3.7 | 5 |
| 49 | Measuring and Interpreting Oxygen Consumption Rates in Whole Fly Head Segments. Journal of Visualized Experiments, 2019, , . | 0.3 | 1 |
| 50 | Distinct metabolic adaptation of liver circadian pathways to acute and chronic patterns of alcohol intake. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 25250-25259. | 7.1 | 38 |
| 51 | Toll-like Receptor Signaling Rewires Macrophage Metabolism and Promotes Histone Acetylation via ATP-Citrate Lyase. Immunity, 2019, 51, 997-1011.e7. | 14.3 | 216 |
| 52 | MIR sequences recruit zinc finger protein ZNF768 to expressed genes. Nucleic Acids Research, 2019, 47, 700-715. | 14.5 | 14 |
| 53 | Shelterin and subtelomeric <scp>DNA</scp> sequences control nucleosome maintenance and genome stability. EMBO Reports, 2019, 20, . | 4.5 | 30 |
| 54 | Analog-sensitive cell line identifies cellular substrates of CDK9. Oncotarget, 2019, 10, 6934-6943. | 1.8 | 18 |

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| 55 | Analysis of Histone Modifications by Mass Spectrometry. Current Protocols in Protein Science, 2018, 92, e54. | 2.8 | 15 |
| 56 | Structural Architecture of the Nucleosome Remodeler ISWI Determined from Cross-Linking, Mass Spectrometry, SAXS, and Modeling. Structure, 2018, 26, 282-294.e6. | 3.3 | 11 |
| 57 | Tyrosine-1 of RNA Polymerase II CTD Controls Global Termination of Gene Transcription in Mammals. Molecular Cell, 2018, 69, 48-61.e6. | 9.7 | 66 |
| 58 | Regulation and function of H3K36 di-methylation by the trithorax-group protein complex AMC. Development (Cambridge), 2018, 145, . | 2.5 | 33 |
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| 61 | Detection of Histone Modification Dynamics during the Cell Cycle by MS-Based Proteomics. Methods in Molecular Biology, 2018, 1832, 61-74. | 0.9 | 2 |
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| 63 | Coronin 1A, a novel player in integrin biology, controls neutrophil trafficking in innate immunity. Blood, 2017, 130, 847-858. | 1.4 | 56 |
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| 65 | Adrenergic Signaling Strengthens Cardiac Myocyte Cohesion. Circulation Research, 2017, 120, 1305-1317. | 4.5 | 55 |
| 66 | NO Augments Endothelial Reactivity by Reducing Myoendothelial Calcium Signal Spreading. Arteriosclerosis, Thrombosis, and Vascular Biology, 2017, 37, 2280-2290. | 2.4 | 18 |
| 67 | Ubiquitome Analysis Reveals PCNA-Associated Factor 15 (PAF15) as a Specific Ubiquitination Target of UHRF1 in Embryonic Stem Cells. Journal of Molecular Biology, 2017, 429, 3814-3824. | 4.2 | 43 |
| 68 | PP32 and SET/TAF-l \hat{l}^2 proteins regulate the acetylation of newly synthesized histone H4. Nucleic Acids Research, 2017, 45, 11700-11710. | 14.5 | 21 |
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| 73 | Life span extension by targeting a link between metabolism and histone acetylation in <i>Drosophila</i> . EMBO Reports, 2016, 17, 455-469. | 4.5 | 116 |
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| 77 | Specific threonine-4 phosphorylation and function of RNA polymerase II CTD during M phase progression. Scientific Reports, 2016, 6, 27401. | 3.3 | 17 |
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| 82 | Combinatorial Histone Acetylation Patterns Are Generated by Motif-Specific Reactions. Cell Systems, 2016, 2, 49-58. | 6.2 | 19 |
| 83 | MALDI imaging mass spectrometry as a novel tool for detecting histone modifications in clinical tissue samples. Expert Review of Proteomics, 2016, 13, 275-284. | 3.0 | 13 |
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| 85 | Assembly of methylated KDM1A and CHD1 drives androgen receptor–dependent transcription and translocation. Nature Structural and Molecular Biology, 2016, 23, 132-139. | 8.2 | 70 |
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| 88 | Identification of Drosophila centromere associated proteins by quantitative affinity purification-mass spectrometry. Data in Brief, 2015, 4, 544-550. | 1.0 | 8 |
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| 90 | DEAD-box helicase DDX27 regulates 3′ end formation of ribosomal 47S RNA and stably associates with the PeBoW-complex. Experimental Cell Research, 2015, 334, 146-159. | 2.6 | 26 |

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| 91 | Two distinct modes for propagation of histone PTMs across the cell cycle. Genes and Development, 2015, 29, 585-590. | 5.9 | 334 |
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| 93 | The Epoxyeicosatrienoic Acid Pathway Enhances Hepatic Insulin Signaling and is Repressed in Insulin-Resistant Mouse Liver*. Molecular and Cellular Proteomics, 2015, 14, 2764-2774. | 3.8 | 13 |
| 94 | Site-specific methylation and acetylation of lysine residues in the C-terminal domain (CTD) of RNA polymerase II. Transcription, 2015, 6, 91-101. | 3.1 | 22 |
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| 100 | Epstein–Barr virus-mediated transformation of B cells induces global chromatin changes independent to the acquisition of proliferation. Nucleic Acids Research, 2014, 42, 249-263. | 14.5 | 34 |
| 101 | Circadian Control of Fatty Acid Elongation by SIRT1 Protein-mediated Deacetylation of Acetyl-coenzyme A Synthetase 1. Journal of Biological Chemistry, 2014, 289, 6091-6097. | 3.4 | 61 |
| 102 | Identification of novel <i>Drosophila</i> centromereâ€associated proteins. Proteomics, 2014, 14, 2167-2178. | 2.2 | 28 |
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| 104 | Ablation of D2 autoreceptors causes epigenetic reprogramming of cortical neurons. Molecular Psychiatry, 2014, 19, 1153-1153. | 7.9 | 3 |
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| 106 | Bioinformatic analysis of proteomics data. BMC Systems Biology, 2014, 8, S3. | 3.0 | 131 |
| 107 | The novel component Kgd4 recruits the E3 subunit to the mitochondrial \hat{l} ±-ketoglutarate dehydrogenase. Molecular Biology of the Cell, 2014, 25, 3342-3349. | 2.1 | 43 |
| 108 | Mislocalization of the Centromeric Histone Variant CenH3/CENP-A in Human Cells Depends on the Chaperone DAXX. Molecular Cell, 2014, 53, 631-644. | 9.7 | 214 |

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| 110 | Stepwise Evolution of Essential Centromere Function in a <i>Drosophila</i> Neogene. Science, 2013, 340, 1211-1214. | 12.6 | 94 |
| 111 | Mouse cytomegalovirus egress protein pM50 interacts with cellular endophilin-A2. Cellular Microbiology, 2013, 15, 335-351. | 2.1 | 23 |
| 112 | Circadian acetylome reveals regulation of mitochondrial metabolic pathways. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3339-3344. | 7.1 | 133 |
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| 116 | Myb-binding Protein 1a (Mybbp1a) Regulates Levels and Processing of Pre-ribosomal RNA. Journal of Biological Chemistry, 2012, 287, 24365-24377. | 3.4 | 37 |
| 117 | Probing the Conformation of the ISWI ATPase Domain With Genetically Encoded Photoreactive Crosslinkers and Mass Spectrometry. Molecular and Cellular Proteomics, 2012, 11, M111.012088. | 3.8 | 45 |
| 118 | Characterization of the insertase for \hat{l}^2 -barrel proteins of the outer mitochondrial membrane. Journal of Cell Biology, 2012, 199, 599-611. | 5.2 | 43 |
| 119 | Developmental regulation of N-terminal H2B methylation in Drosophila melanogaster. Nucleic Acids Research, 2012, 40, 1536-1549. | 14.5 | 28 |
| 120 | Role of the AAA protease Yme1 in folding of proteins in the intermembrane space of mitochondria. Molecular Biology of the Cell, 2012, 23, 4335-4346. | 2.1 | 50 |
| 121 | Connecting Threads: Epigenetics and Metabolism. Cell, 2012, 148, 24-28. | 28.9 | 282 |
| 122 | MSL2 Combines Sensor and Effector Functions in Homeostatic Control of the Drosophila Dosage Compensation Machinery. Molecular Cell, 2012, 48, 647-654. | 9.7 | 31 |
| 123 | Df31 Protein and snoRNAs Maintain Accessible Higher-Order Structures of Chromatin. Molecular Cell, 2012, 48, 434-444. | 9.7 | 108 |
| 124 | Impairment of prostate cancer cell growth by a selective and reversible lysineâ€specific demethylase 1 inhibitor. International Journal of Cancer, 2012, 131, 2704-2709. | 5.1 | 118 |
| 125 | Secretome protein enrichment identifies physiological BACE1 protease substrates in neurons. EMBO Journal, 2012, 31, 3157-3168. | 7.8 | 279 |
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| 127 | The RNA Helicase Rm62 Cooperates with SU(VAR)3-9 to Re-Silence Active Transcription in Drosophila melanogaster. PLoS ONE, 2011, 6, e20761. | 2.5 | 9 |
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| 134 | Fast signals and slow marks: the dynamics of histone modifications. Trends in Biochemical Sciences, 2010, 35, 618-626. | 7.5 | 268 |
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| 138 | Replication Stress Interferes with Histone Recycling and Predeposition Marking of New Histones. Molecular Cell, 2010, 37, 736-743. | 9.7 | 242 |
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| 144 | The HP1α–CAF1–SetDB1â€containing complex provides H3K9me1 for Suv39â€mediated K9me3 in pericentine heterochromatin. EMBO Reports, 2009, 10, 769-775. | ric 4.5 | 201 |

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| 146 | Dynamics of Adaptive Microevolution of Hypermutable <i>Pseudomonas aeruginosa </i> Pulmonary Infection in Patients with Cystic Fibrosis. Journal of Infectious Diseases, 2009, 200, 118-130. | 4.0 | 155 |
| 147 | Drosophila HP1c Is Regulated by an Auto-Regulatory Feedback Loop through Its Binding Partner Woc. PLoS ONE, 2009, 4, e5089. | 2.5 | 21 |
| 148 | ESC, ESCL and their roles in Polycomb Group mechanisms. Mechanisms of Development, 2008, 125, 527-541. | 1.7 | 34 |
| 149 | Analysis of Histone Modifications by Mass Spectrometry. Current Protocols in Protein Science, 2008, 51, Unit 14.10. | 2.8 | 32 |
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| 154 | HP1 Binding to Chromatin Methylated at H3K9 Is Enhanced by Auxiliary Factors. Molecular and Cellular Biology, 2007, 27, 453-465. | 2.3 | 115 |
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| 156 | Activation of RNA Polymerase I Transcription by Cockayne Syndrome Group B Protein and Histone Methyltransferase G9a. Molecular Cell, 2007, 27, 585-595. | 9.7 | 147 |
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| 159 | Site-specific acetylation of ISWI by GCN5. BMC Molecular Biology, 2007, 8, 73. | 3.0 | 40 |
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| 164 | Proteome analysis of mitochondrial outer membrane from Neurospora crassa. Proteomics, 2006, 6, 72-80. | 2.2 | 74 |
| 165 | A \hat{I}^3 -secretase-like intramembrane cleavage of TNF \hat{I}^\pm by the GxGD aspartyl protease SPPL2b. Nature Cell Biology, 2006, 8, 894-896. | 10.3 | 130 |
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