

List of Publications by Year in descending order

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35
papers

36,812
citations

218381

26
h-index

360668

35
g-index

37
all docs

37
docs citations

37
times ranked

40067
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | The Protein Data Bank. <i>Nucleic Acids Research</i> , 2000, 28, 235-242. | 6.5 | 31,087 |
| 2 | RCSB Protein Data Bank: biological macromolecular structures enabling research and education in fundamental biology, biomedicine, biotechnology and energy. <i>Nucleic Acids Research</i> , 2019, 47, D464-D474. | 6.5 | 918 |
| 3 | Protein Data Bank: the single global archive for 3D macromolecular structure data. <i>Nucleic Acids Research</i> , 2019, 47, D520-D528. | 6.5 | 671 |
| 4 | The Protein Data Bank and the challenge of structural genomics. <i>Nature Structural Biology</i> , 2000, 7, 957-959. | 9.7 | 511 |
| 5 | The RCSB PDB information portal for structural genomics. <i>Nucleic Acids Research</i> , 2006, 34, D302-D305. | 6.5 | 334 |
| 6 | The Protein Data Bank and structural genomics. <i>Nucleic Acids Research</i> , 2003, 31, 489-491. | 6.5 | 331 |
| 7 | The RCSB Protein Data Bank: a redesigned query system and relational database based on the mmCIF schema. <i>Nucleic Acids Research</i> , 2004, 33, D233-D237. | 6.5 | 303 |
| 8 | The Protein Data Bank: unifying the archive. <i>Nucleic Acids Research</i> , 2002, 30, 245-248. | 6.5 | 261 |
| 9 | Tools for the automatic identification and classification of RNA base pairs. <i>Nucleic Acids Research</i> , 2003, 31, 3450-3460. | 6.5 | 240 |
| 10 | RCSB Protein Data Bank: Sustaining a living digital data resource that enables breakthroughs in scientific research and biomedical education. <i>Protein Science</i> , 2018, 27, 316-330. | 3.1 | 219 |
| 11 | RNA backbone: Consensus all-angle conformers and modular string nomenclature (an RNA Ontology) Tj ETQq1 1 0,784314 rgBT /Ovele | 1.6 | 216 |
| 12 | 2017 publication guidelines for structural modelling of small-angle scattering data from biomolecules in solution: an update. <i>Acta Crystallographica Section D: Structural Biology</i> , 2017, 73, 710-728. | 1.1 | 205 |
| 13 | The Nucleic Acid Database: new features and capabilities. <i>Nucleic Acids Research</i> , 2014, 42, D114-D122. | 6.5 | 194 |
| 14 | Geometric Parameters in Nucleic Acids: \hat{A} Nitrogenous Bases. <i>Journal of the American Chemical Society</i> , 1996, 118, 509-518. | 6.6 | 191 |
| 15 | PDBML: the representation of archival macromolecular structure data in XML. <i>Bioinformatics</i> , 2005, 21, 988-992. | 1.8 | 154 |
| 16 | Remediation of the protein data bank archive. <i>Nucleic Acids Research</i> , 2007, 36, D426-D433. | 6.5 | 136 |
| 17 | The distribution and query systems of the RCSB Protein Data Bank. <i>Nucleic Acids Research</i> , 2004, 32, 223D-225. | 6.5 | 108 |
| 18 | RNAML: A standard syntax for exchanging RNA information. <i>Rna</i> , 2002, 8, 707-717. | 1.6 | 91 |

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|----|--|-----|-----------|
| 19 | Report of the wwPDB Small-Angle Scattering Task Force: Data Requirements for Biomolecular Modeling and the PDB. <i>Structure</i> , 2013, 21, 875-881. | 1.6 | 77 |
| 20 | The Nucleic Acid Database. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2002, 58, 889-898. | 2.5 | 57 |
| 21 | Announcing mandatory submission of PDBx/mmCIF format files for crystallographic depositions to the Protein Data Bank (PDB). <i>Acta Crystallographica Section D: Structural Biology</i> , 2019, 75, 451-454. | 1.1 | 46 |
| 22 | Validation of Protein Structures for Protein Data Bank. <i>Methods in Enzymology</i> , 2003, 374, 370-385. | 0.4 | 43 |
| 23 | A framework for scientific data modeling and automated software development. <i>Bioinformatics</i> , 2005, 21, 1678-1684. | 1.8 | 42 |
| 24 | NMR Exchange Format: a unified and open standard for representation of NMR restraint data. <i>Nature Structural and Molecular Biology</i> , 2015, 22, 433-434. | 3.6 | 40 |
| 25 | Improving the representation of peptide-like inhibitor and antibiotic molecules in the Protein Data Bank. <i>Biopolymers</i> , 2014, 101, 659-668. | 1.2 | 31 |
| 26 | The Protein Data Bank. , 2003, , 389-405. | | 29 |
| 27 | Crystallography and Databases. <i>Data Science Journal</i> , 2017, 16, . | 0.6 | 28 |
| 28 | Design of a data model for developing laboratory information management and analysis systems for protein production. <i>Proteins: Structure, Function and Bioinformatics</i> , 2004, 58, 278-284. | 1.5 | 27 |
| 29 | Archiving and disseminating integrative structure models. <i>Journal of Biomolecular NMR</i> , 2019, 73, 385-398. | 1.6 | 20 |
| 30 | Extension of the sasCIF format and its applications for data processing and deposition. <i>Journal of Applied Crystallography</i> , 2016, 49, 302-310. | 1.9 | 18 |
| 31 | Realism about PDB. <i>Nature Biotechnology</i> , 2007, 25, 845-846. | 9.4 | 17 |
| 32 | RCSB Protein Data Bank 1D tools and services. <i>Bioinformatics</i> , 2021, 36, 5526-5527. | 1.8 | 15 |
| 33 | Chemical annotation of small and peptide-like molecules at the Protein Data Bank. <i>Database: the Journal of Biological Databases and Curation</i> , 2013, 2013, bat079. | 1.4 | 14 |
| 34 | The Nucleic Acid Database: A Resource for Nucleic Acid Science. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 1998, 54, 1095-1104. | 2.5 | 13 |
| 35 | The evolution of the RCSB Protein Data Bank website. <i>Wiley Interdisciplinary Reviews: Computational Molecular Science</i> , 2011, 1, 782-789. | 6.2 | 7 |