

# Lora Mak

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9049020/publications.pdf>

Version: 2024-02-01

10  
papers

1,442  
citations

1040056

9  
h-index

1474206

9  
g-index

10  
all docs

10  
docs citations

10  
times ranked

2446  
citing authors

#	ARTICLE	IF	CITATIONS
1	Protein Data Bank: the single global archive for 3D macromolecular structure data. <i>Nucleic Acids Research</i> , 2019, 47, D520-D528.	14.5	671
2	Validation of Structures in the Protein Data Bank. <i>Structure</i> , 2017, 25, 1916-1927.	3.3	210
3	PDBe: improved accessibility of macromolecular structure data from PDB and EMDB. <i>Nucleic Acids Research</i> , 2016, 44, D385-D395.	14.5	131
4	OneDep: Unified wwPDB System for Deposition, Biocuration, and Validation of Macromolecular Structures in the PDB Archive. <i>Structure</i> , 2017, 25, 536-545.	3.3	130
5	PDBe: improved findability of macromolecular structure data in the PDB. <i>Nucleic Acids Research</i> , 2020, 48, D335-D343.	14.5	86
6	PDBe: towards reusable data delivery infrastructure at protein data bank in Europe. <i>Nucleic Acids Research</i> , 2018, 46, D486-D492.	14.5	76
7	Metabase: a cheminformatics and bioinformatics database for small molecule transporter data analysis and (Q)SAR modeling. <i>Journal of Cheminformatics</i> , 2015, 7, 31.	6.1	47
8	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.	2.3	46
9	Worldwide Protein Data Bank biocuration supporting open access to high-quality 3D structural biology data. <i>Database: the Journal of Biological Databases and Curation</i> , 2018, 2018, .	3.0	45
10	Chapter 7. Methods and Resources for Transport Proteins in <i>Bioinformatics and Cheminformatics</i> . , 0, 195-226.		0