

# Andrey Lebedev

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8263679/publications.pdf>

Version: 2024-02-01

8

papers

628

citations

1478505

6

h-index

1720034

7

g-index

8

all docs

8

docs citations

8

times ranked

1499

citing authors

#	ARTICLE	IF	CITATIONS
1	Modelling covalent linkages in <i>CCP</i>4. <i>Acta Crystallographica Section D: Structural Biology</i> , 2021, 77, 712-726.	2.3	10
2	Multitasking in the gut: the X-ray structure of the multidomain BbgIII from <i>Bifidobacterium bifidum</i> offers possible explanations for its alternative functions. <i>Acta Crystallographica Section D: Structural Biology</i> , 2021, 77, 1564-1578.	2.3	0
3	Using <i>Phaser</i> and ensembles to improve the performance of <i>SIMBAD</i>. <i>Acta Crystallographica Section D: Structural Biology</i> , 2020, 76, 1-8.	2.3	7
4	<i>CCP</i>4<i>i</i>2: the new graphical user interface to the <i>CCP</i>4 program suite. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 68-84.	2.3	382
5	Pathological macromolecular crystallographic data affected by twinning, partial-disorder and exhibiting multiple lattices for testing of data processing and refinement tools. <i>Scientific Reports</i> , 2018, 8, 14876.	3.3	11
6	Distributed computing for macromolecular crystallography. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 143-151.	2.3	54
7	<i>SIMBAD</i>: a sequence-independent molecular-replacement pipeline. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 595-605.	2.3	25
8	<i>MoRDa</i>, an automatic molecular replacement pipeline. <i>Acta Crystallographica Section A: Foundations and Advances</i> , 2015, 71, s19-s19.	0.1	139