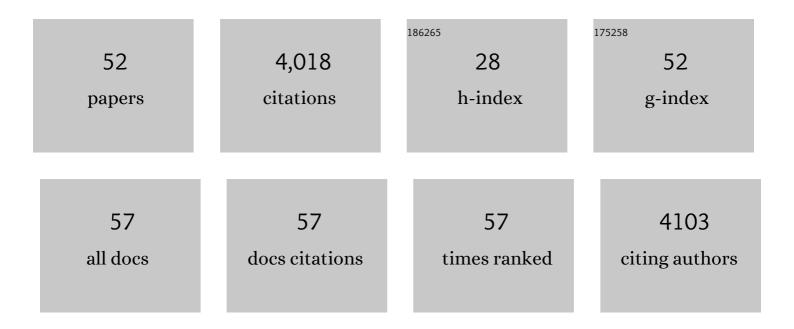
Oleksandr Yefanov

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Unsupervised learning approaches to characterizing heterogeneous samples using X-ray single-particle imaging. IUCrJ, 2022, 9, 204-214.	2.2	9
2	Using diffraction losses of X-rays in a single crystal for determination of its lattice parameters as well as for monochromator calibration. Journal of Synchrotron Radiation, 2022, 29, 369-376.	2.4	2
3	Robust ptychographic X-ray speckle tracking with multilayer Laue lenses. Optics Express, 2022, 30, 25450.	3.4	1
4	Ultrafast structural changes within a photosynthetic reaction centre. Nature, 2021, 589, 310-314.	27.8	47
5	Synchronous RNA conformational changes trigger ordered phase transitions in crystals. Nature Communications, 2021, 12, 1762.	12.8	17
6	Determination of the Exact Orientation of Single-Crystal X-ray Optics from Its Glitch Spectrum and Modeling of Glitches for an Arbitrary Configuration. Crystals, 2021, 11, 504.	2.2	5
7	Data reduction for serial crystallography using a robust peak finder. Journal of Applied Crystallography, 2021, 54, 1360-1378.	4.5	10
8	Observation of substrate diffusion and ligand binding in enzyme crystals using high-repetition-rate mix-and-inject serial crystallography. IUCrJ, 2021, 8, 878-895.	2.2	44
9	Pink-beam serial femtosecond crystallography for accurate structure-factor determination at an X-ray free-electron laser. IUCrJ, 2021, 8, 905-920.	2.2	11
10	3D diffractive imaging of nanoparticle ensembles using an x-ray laser. Optica, 2021, 8, 15.	9.3	48
11	Suppressing Diffraction-Related Intensity Losses in Transmissive Single-Crystal X-ray Optics. Crystals, 2021, 11, 1561.	2.2	3
12	Time-resolved serial femtosecond crystallography at the European XFEL. Nature Methods, 2020, 17, 73-78.	19.0	110
13	Serial protein crystallography in an electron microscope. Nature Communications, 2020, 11, 996.	12.8	69
14	Ultracompact 3D microfluidics for time-resolved structural biology. Nature Communications, 2020, 11, 657.	12.8	106
15	Predicting glitches of intensity in single-crystal diamond CRLs. AIP Conference Proceedings, 2020, , .	0.4	4
16	Ptychographic X-ray speckle tracking with multi-layer Laue lens systems. Journal of Applied Crystallography, 2020, 53, 927-936.	4.5	11
17	<i>pinkIndexer</i> – a universal indexer for pink-beam X-ray and electron diffraction snapshots. Acta Crystallographica Section A: Foundations and Advances, 2020, 76, 121-131.	0.1	28
18	Evaluation of serial crystallographic structure determination within megahertz pulse trains. Structural Dynamics, 2019, 6, 064702.	2.3	26

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19	On-chip crystallization for serial crystallography experiments and on-chip ligand-binding studies. IUCrJ, 2019, 6, 714-728.	2.2	41
20	1 kHz fixed-target serial crystallography using a multilayer monochromator and an integrating pixel detector. IUCrJ, 2019, 6, 927-937.	2.2	35
21	<i>XGANDALF</i> – extended gradient descent algorithm for lattice finding. Acta Crystallographica Section A: Foundations and Advances, 2019, 75, 694-704.	0.1	80
22	Femtosecond X-ray diffraction from an aerosolized beam of protein nanocrystals. Journal of Applied Crystallography, 2018, 51, 133-139.	4.5	18
23	Megahertz serial crystallography. Nature Communications, 2018, 9, 4025.	12.8	147
24	X-ray focusing with efficient high-NA multilayer Laue lenses. Light: Science and Applications, 2018, 7, 17162-17162.	16.6	114
25	Enzyme intermediates captured "on the fly―by mix-and-inject serial crystallography. BMC Biology, 2018, 16, 59.	3.8	117
26	Rapid sample delivery for megahertz serial crystallography at X-ray FELs. IUCrJ, 2018, 5, 574-584.	2.2	52
27	Structural enzymology using X-ray free electron lasers. Structural Dynamics, 2017, 4, 044003.	2.3	92
28	Atomic structure of granulin determined from native nanocrystalline granulovirus using an X-ray free-electron laser. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 2247-2252.	7.1	65
29	Double-flow focused liquid injector for efficient serial femtosecond crystallography. Scientific Reports, 2017, 7, 44628.	3.3	90
30	From Macrocrystals to Microcrystals: A Strategy for Membrane Protein Serial Crystallography. Structure, 2017, 25, 1461-1468.e2.	3.3	21
31	Probing Dynamics in Colloidal Crystals with Pump-Probe Experiments at LCLS: Methodology and Analysis. Applied Sciences (Switzerland), 2017, 7, 519.	2.5	4
32	Post-sample aperture for low background diffraction experiments at X-ray free-electron lasers. Journal of Synchrotron Radiation, 2017, 24, 1296-1298.	2.4	8
33	FELIX: an algorithm for indexing multiple crystallites in X-ray free-electron laser snapshot diffraction images. Journal of Applied Crystallography, 2017, 50, 1075-1083.	4.5	27
34	Atomic structure of granulin determined from native nanocrystalline granulovirus using an X-ray free-electron laser. Acta Crystallographica Section A: Foundations and Advances, 2017, 73, a292-a293.	0.1	2
35	Mix-and-diffuse serial synchrotron crystallography. IUCrJ, 2017, 4, 769-777.	2.2	98
36	Analysis of XFEL serial diffraction data from individual crystalline fibrils. IUCrJ, 2017, 4, 795-811.	2.2	16

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37	Femtosecond structural dynamics drives the trans/cis isomerization in photoactive yellow protein. Science, 2016, 352, 725-729.	12.6	348
38	X-ray laser diffraction for structure determination of the rhodopsin-arrestin complex. Scientific Data, 2016, 3, 160021.	5.3	51
39	Recent developments in <i>CrystFEL</i> . Journal of Applied Crystallography, 2016, 49, 680-689.	4.5	222
40	Perspectives for imaging single protein molecules with the present design of the European XFEL. Structural Dynamics, 2015, 2, 041702.	2.3	29
41	Accurate determination of segmented X-ray detector geometry. Optics Express, 2015, 23, 28459.	3.4	69
42	Trace phase detection and strain characterization from serial X-ray free-electron laser crystallography of a Pr _{0.5} Ca _{0.5} MnO ₃ powder. Powder Diffraction, 2015, 30, S25-S30.	0.2	1
43	Ternary structure reveals mechanism of a membrane diacylglycerol kinase. Nature Communications, 2015, 6, 10140.	12.8	30
44	Structural Evolution of Colloidal Crystal Films in the Process of Melting Revealed by Bragg Peak Analysis. Langmuir, 2015, 31, 5274-5283.	3.5	27
45	Crystal structure of rhodopsin bound to arrestin by femtosecond X-ray laser. Nature, 2015, 523, 561-567.	27.8	683
46	High numerical aperture multilayer Laue lenses. Scientific Reports, 2015, 5, 9892.	3.3	89
47	Time-resolved serial crystallography captures high-resolution intermediates of photoactive yellow protein. Science, 2014, 346, 1242-1246.	12.6	418
48	Mapping the continuous reciprocal space intensity distribution of X-ray serial crystallography. Philosophical Transactions of the Royal Society B: Biological Sciences, 2014, 369, 20130333.	4.0	29
49	Serial crystallography on <i>in vivo</i> grown microcrystals using synchrotron radiation. IUCrJ, 2014, 1, 87-94.	2.2	204
50	Room-temperature macromolecular serial crystallography using synchrotron radiation. IUCrJ, 2014, 1, 204-212.	2.2	221
51	Accessible reciprocal-space region for non-coplanar Bragg and Laue geometries. Journal of Applied Crystallography, 2008, 41, 110-114.	4.5	5
52	<i>XVis</i> : an educational open-source program for demonstration of reciprocal-space construction and diffraction principles. Journal of Applied Crystallography, 2008, 41, 647-652.	4.5	6