Victor A Streltsov

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Structure of the insulin receptor ectodomain reveals a folded-over conformation. Nature, 2006, 443, 218-221.	27.8	277
2	Structure of the Haemagglutinin-neuraminidase from Human Parainfluenza Virus Type III. Journal of Molecular Biology, 2004, 335, 1343-1357.	4.2	200
3	Multipole analysis of the electron density in triphylite, LiFePO4, using X-ray diffraction data. Acta Crystallographica Section B: Structural Science, 1993, 49, 147-153.	1.8	196
4	Mechanism-Based Covalent Neuraminidase Inhibitors with Broad-Spectrum Influenza Antiviral Activity. Science, 2013, 340, 71-75.	12.6	175
5	Electron density and optical anisotropy in rhombohedral carbonates. III. Synchrotron X-ray studies of CaCO3, MgCO3 and MnCO3. Acta Crystallographica Section B: Structural Science, 1995, 51, 929-939.	1.8	155
6	X-ray study of the electron density in calcite, CaCo3. Acta Crystallographica Section B: Structural Science, 1993, 49, 636-641.	1.8	148
7	Synchrotron X-ray study of the electron density in α-Fe2O3. Acta Crystallographica Section B: Structural Science, 1994, 50, 435-441.	1.8	140
8	The insulin and EGF receptor structures: new insights into ligand-induced receptor activation. Trends in Biochemical Sciences, 2007, 32, 129-137.	7.5	122
9	Structural evidence for evolution of shark Ig new antigen receptor variable domain antibodies from a cell-surface receptor. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 12444-12449.	7.1	119
10	The Structure of the Amyloid-β Peptide High-Affinity Copper II Binding Site in Alzheimer Disease. Biophysical Journal, 2008, 95, 3447-3456.	0.5	108
11	Iron, Copper, and Zinc Concentration in Aβ Plaques in the APP/PS1 Mouse Model of Alzheimer's Disease Correlates with Metal Levels in the Surrounding Neuropil. ACS Chemical Neuroscience, 2017, 8, 629-637.	3.5	107
12	Structure of an IgNAR-AMA1 Complex: Targeting a Conserved Hydrophobic Cleft Broadens Malarial Strain Recognition. Structure, 2007, 15, 1452-1466.	3.3	101
13	Crystal Structure of the Amyloid-β p3 Fragment Provides a Model for Oligomer Formation in Alzheimer's Disease. Journal of Neuroscience, 2011, 31, 1419-1426.	3.6	99
14	Ammonium hydroxide treatment of Aβ produces an aggregate free solution suitable for biophysical and cell culture characterization. PeerJ, 2013, 1, e73.	2.0	93
15	Naturally occurring polyphenolic inhibitors of amyloid beta aggregation. Bioorganic and Medicinal Chemistry Letters, 2014, 24, 3108-3112.	2.2	76
16	Structural and Functional Basis of Resistance to Neuraminidase Inhibitors of Influenza B Viruses. Journal of Medicinal Chemistry, 2010, 53, 6421-6431.	6.4	75
17	Structure of a shark IgNAR antibody variable domain and modeling of an early-developmental isotype. Protein Science, 2005, 14, 2901-2909.	7.6	68
18	A synchrotron X-ray study of the electron density in YFeO3. Acta Crystallographica Section B: Structural Science, 1995, 51, 921-929.	1.8	65

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19	A synchrotron X-ray study of the electron density in C-type rare earth oxides. Acta Crystallographica Section B: Structural Science, 1996, 52, 414-422.	1.8	64
20	Hematinâ~'Hematin Self-Association States Involved in the Formation and Reactivity of the Malaria Parasite Pigment, Hemozoin. Biochemistry, 2010, 49, 6804-6811.	2.5	57
21	A synchrotron X-ray study of the electron density in SmFeO3. Acta Crystallographica Section B: Structural Science, 1996, 52, 406-413.	1.8	52
22	Structural insights into ligandâ€induced activation of the insulin receptor. Acta Physiologica, 2008, 192, 3-9.	3.8	50
23	L22 Ribosomal Protein and Effect of Its Mutation on Ribosome Resistance to Erythromycin. Journal of Molecular Biology, 2002, 322, 635-644.	4.2	48
24	Haloalkane Dehalogenase LinB fromSphingomonas paucimobilisUT26:Â X-ray Crystallographic Studies of Dehalogenation of Brominated Substratesâ€,‡. Biochemistry, 2003, 42, 10104-10112.	2.5	43
25	Dimerisation strategies for shark IgNAR single domain antibody fragments. Journal of Immunological Methods, 2006, 315, 171-184.	1.4	43
26	Structural Rationale for Low-Nanomolar Binding of Transition State Mimics to a Family GH3 β-d-Glucan Glucohydrolase from Barleyâ€,‡. Biochemistry, 2005, 44, 16529-16539.	2.5	42
27	Inhibition of amyloid beta-induced synaptotoxicity by a pentapeptide derived from the glycine zipper region of the neurotoxic peptide. Neurobiology of Aging, 2013, 34, 2805-2814.	3.1	41
28	Synchrotron X-ray and ab initio studies of Î ² -Si3N4. Acta Crystallographica Section B: Structural Science, 2004, 60, 388-405.	1.8	38
29	Structural insights into the interaction of platinum-based inhibitors with the Alzheimer's disease amyloid-l ² peptide. Chemical Communications, 2013, 49, 11364.	4.1	38
30	Synchrotron X-ray study of the electron density in RFeO3 (R = Nd, Dy). Acta Crystallographica Section B: Structural Science, 1999, 55, 1-7.	1.8	35
31	Differential Receptor Binding and Regulatory Mechanisms for the Lymphangiogenic Growth Factors Vascular Endothelial Growth Factor (VEGF)-C and -D. Journal of Biological Chemistry, 2016, 291, 27265-27278.	3.4	35
32	In vitro improvement of a shark IgNAR antibody by Qβ replicase mutation and ribosome display mimics in vivo affinity maturation. Immunology Letters, 2006, 107, 163-168.	2.5	34
33	Lysine postâ€translational modification of glyceraldehydeâ€3â€phosphate dehydrogenase regulates hepatic and systemic metabolism. FASEB Journal, 2017, 31, 2592-2602.	0.5	31
34	Shark IgNAR antibody mimotopes target a murine immunoglobulin through extended CDR3 loop structures. Proteins: Structure, Function and Bioinformatics, 2008, 71, 119-130.	2.6	27
35	Substrate mediated reduction of copper-amyloid-β complex in Alzheimer's disease. Chemical Communications, 2008, , 3169.	4.1	27
36	In vitro passaging of a pandemic H1N1/09 virus selects for viruses with neuraminidase mutations conferring high-level resistance to oseltamivir and peramivir, but not to zanamivir. Journal of Antimicrobial Chemotherapy, 2012, 67, 1874-1883.	3.0	27

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37	Synchrotron X-ray analysis of the electron density in CoF2 and ZnF2. Acta Crystallographica Section B: Structural Science, 2001, 57, 128-135.	1.8	26
38	Germline humanization of a murine Al̂² antibody and crystal structure of the humanized recombinant Fab fragment. Protein Science, 2010, 19, 299-308.	7.6	25
39	Structural Insight into Redox Dynamics of Copper Bound N-Truncated Amyloid-β Peptides from <i>in Situ</i> X-ray Absorption Spectroscopy. Inorganic Chemistry, 2018, 57, 11422-11435.	4.0	25
40	Charge density analysis from complementary high energy synchrotron X-ray and electron diffraction data. Journal of Physics and Chemistry of Solids, 2001, 62, 2109-2117.	4.0	23
41	X-ray absorption and diffraction studies of the metal binding sites in amyloid β-peptide. European Biophysics Journal, 2008, 37, 257-263.	2.2	23
42	Structural studies of the tethered N-terminus of the Alzheimer's disease amyloid-β peptide. Proteins: Structure, Function and Bioinformatics, 2013, 81, 1748-1758.	2.6	22
43	X-ray laser–induced electron dynamics observed by femtosecond diffraction from nanocrystals of Buckminsterfullerene. Science Advances, 2016, 2, e1601186.	10.3	20
44	Discovery of processive catalysis by an exo-hydrolase with a pocket-shaped active site. Nature Communications, 2019, 10, 2222.	12.8	20
45	Alkoxide coordination of iron(iii) protoporphyrin IX by antimalarial quinoline methanols: a key interaction observed in the solid-state and solution. Dalton Transactions, 2015, 44, 16767-16777.	3.3	19
46	Continuous X-ray diffractive field in protein nanocrystallography. Acta Crystallographica Section A: Foundations and Advances, 2013, 69, 108-118.	0.3	18
47	Synchrotron X-ray study of Er3Al5O12 and Yb3Al5O12 garnets. Acta Crystallographica Section B: Structural Science, 2001, 57, 136-141.	1.8	17
48	A Combination Method of Charge Density Measurement in Hard Materials Using Accurate, Quantitative Electron and X-ray Diffraction: The α-Al2O3 Case. Microscopy and Microanalysis, 2003, 9, 419-427.	0.4	17
49	Domain I of ribosomal protein L1 is sufficient for specific RNA binding. Nucleic Acids Research, 2007, 35, 7389-7395.	14.5	17
50	Unprecedented conformational flexibility revealed in the ligand-binding domains of the <i>Bovicola ovis</i> ecdysone receptor (EcR) and ultraspiracle (USP) subunits. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 1954-1964.	2.5	17
51	X-ray study of the electron density in magnesite MgCO3. Acta Crystallographica Section B: Structural Science, 1993, 49, 980-984.	1.8	16
52	Construction, crystal structure and application of a recombinant protein that lacks the collagenâ€like region of BclA from <i>Bacillus anthracis</i> spores. Biotechnology and Bioengineering, 2008, 99, 774-782.	3.3	15
53	The VD1 Neutralizing Antibody to Vascular Endothelial Growth Factor-D: Binding Epitope and Relationship to Receptor Binding. Journal of Molecular Biology, 2011, 407, 581-593.	4.2	15
54	Preparation of human vascular endothelial growth factor-D for structural and preclinical therapeutic studies. Protein Expression and Purification, 2012, 82, 232-239.	1.3	15

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55	Structure and Function of Ecdysone Receptors—Interactions with Ecdysteroids and Synthetic Agonists. Advances in Insect Physiology, 2012, 43, 299-351.	2.7	15
56	Molecular Structures and Solvation of Free Monomeric and Dimeric Ferriheme in Aqueous Solution: Insights from Molecular Dynamics Simulations and Extended X-ray Absorption Fine Structure Spectroscopy. Inorganic Chemistry, 2014, 53, 10811-10824.	4.0	15
57	Catalytic mechanism and novel receptor binding sites of human parainfluenza virus type 3 hemagglutinin-neuraminidase (hPIV3 HN). Antiviral Research, 2015, 123, 216-223.	4.1	15
58	Synchrotron X-ray electron density in the layered LaOCl structure. Acta Crystallographica Section B: Structural Science, 1996, 52, 576-579.	1.8	14
59	Solution structures of chloroquine–ferriheme complexes modeled using MD simulation and investigated by EXAFS spectroscopy. Journal of Inorganic Biochemistry, 2016, 154, 114-125.	3.5	14
60	Nanocrystallography measurements of early stage synthetic malaria pigment. Journal of Applied Crystallography, 2017, 50, 1533-1540.	4.5	11
61	Synchrotron X-ray analysis of RbTiOAsO4. Acta Crystallographica Section B: Structural Science, 2000, 56, 785-792.	1.8	10
62	Crystal structure study of a β′-copper vanadium bronze, Cu x V2O5 (x = 0.63), by X-ray and convergent beam electron diffraction. Acta Crystallographica Section B: Structural Science, 2005, 61, 17-24.	1.8	10
63	Citrullination of Amyloid-β Peptides in Alzheimer's Disease. ACS Chemical Neuroscience, 2021, 12, 3719-3732.	3.5	10
64	Structure of and electron density in RbTiOAsO4 at 9.6â€K. Acta Crystallographica Section B: Structural Science, 1999, 55, 712-720.	1.8	9
65	Cation movement and phase transitions in KTP isostructures; X-ray study of sodium-doped KTP at 10.5â€K. Acta Crystallographica Section B: Structural Science, 2003, 59, 353-360.	1.8	9
66	Structural and Functional Analysis of Anti-Influenza Activity of 4-, 7-, 8- and 9-Deoxygenated 2,3-Difluoro- <i>N</i> -acetylneuraminic Acid Derivatives. Journal of Medicinal Chemistry, 2018, 61, 1921-1933.	6.4	9
67	Electron density in the sodium vanadium oxide bronze β-Na x V2O5 at 9â€K. Acta Crystallographica Section B: Structural Science, 2001, 57, 244-250.	1.8	8
68	Do sharks have a new antibody lineage?. Immunology Letters, 2005, 97, 159-160.	2.5	8
69	Synchrotron X-ray Imaging of the Electron Density in RFeO3 (R = Y, Ho) Using an APD Detector. Journal of Synchrotron Radiation, 1998, 5, 1309-1316.	2.4	7
70	Structure of an Influenza A virus N9 neuraminidase with a tetrabrachion-domain stalk. Acta Crystallographica Section F, Structural Biology Communications, 2019, 75, 89-97.	0.8	7
71	Isolation, kinetic analysis, and structural characterization of an antibody targeting the <i>Bacillus anthracis</i> major spore surface protein BclA. Proteins: Structure, Function and Bioinformatics, 2011, 79, 1306-1317.	2.6	6
72	A library of AuNPs modified by RAFT polymers of different charge and chain length: high throughput synthesis and synchrotron XFM imaging using a zebrafish larvae model. RSC Advances, 2016, 6, 23550-23563.	3.6	6

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73	Crystallisation of Wild-Type and Variant Forms of a Recombinant Plant Enzyme β-D-Glucan Glucohydrolase from Barley (Hordeum vulgare L.) and Preliminary X-ray Analysis. International Journal of Molecular Sciences, 2010, 11, 2759-2769.	4.1	5
74	Oligomerization and toxicity of $A\hat{l}^2$ fusion proteins. Biochemical and Biophysical Research Communications, 2011, 409, 477-482.	2.1	5
75	Passaging of an influenza A(H1N1)pdm09 virus in a difluoro sialic acid inhibitor selects for a novel, but unfit I106M neuraminidase mutant. Antiviral Research, 2019, 169, 104542.	4.1	5
76	Synchrotron X-ray analysis of the electron density in HoFe2. Acta Crystallographica Section B: Structural Science, 1999, 55, 321-326.	1.8	4
77	Dopant positions in strontium/chromium- and barium-doped KTP, determined with synchrotron X-radiation. Acta Crystallographica Section B: Structural Science, 2000, 56, 980-987.	1.8	4
78	Whole-pattern fitting technique in serial femtosecond nanocrystallography. IUCrJ, 2016, 3, 127-138.	2.2	4
79	Modelling Copper Binding to the Amyloid-β Peptide in Alzheimer. Australian Journal of Chemistry, 2010, 63, 345.	0.9	3
80	Measurements of Long-range Electronic Correlations During Femtosecond Diffraction Experiments Performed on Nanocrystals of Buckminsterfullerene. Journal of Visualized Experiments, 2017, , .	0.3	3
81	Electron density distribution in 3d-metal sesquioxides. Physica Status Solidi A, 1989, 115, 515-521.	1.7	2
82	Redox state and photoreduction control using X-ray spectroelectrochemical techniques – advances in design and fabrication through additive engineering. Journal of Synchrotron Radiation, 2021, 28, 472-479.	2.4	2
83	Defects in Crystals under Pressure. Physica Status Solidi A, 1985, 91, 89-98.	1.7	1
84	Crystals of a mutant form of ribosomal protein L22 rendering bacterial ribosomes resistant to erythromycin. Acta Crystallographica Section D: Biological Crystallography, 2001, 57, 1150-1152.	2.5	1
85	Substitutions at H134 and in the 430-loop region in influenza B neuraminidases can confer reduced susceptibility to multiple neuraminidase inhibitors. Antiviral Research, 2020, 182, 104895.	4.1	1
86	Design of non-aggregating variants of AÎ ² peptide. Biochemical and Biophysical Research Communications, 2014, 453, 449-454.	2.1	0