Kamel El Omari

List of Publications by Year in descending order

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279701 330025 1,681 37 23 37 citations h-index g-index papers 39 39 39 2778 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Crystal structure of the RNA-dependent RNA polymerase from influenza C virus. Nature, 2015, 527, 114-117.	13.7	145
2	Structure of a Pestivirus Envelope Glycoprotein E2 Clarifies Its Role in Cell Entry. Cell Reports, 2013, 3, 30-35.	2.9	124
3	Importance of potassium ions for ribosome structure and function revealed by long-wavelength X-ray diffraction. Nature Communications, 2019, 10, 2519.	5.8	124
4	Influenza Polymerase Can Adopt an Alternative Configuration Involving a Radical Repacking of PB2 Domains. Molecular Cell, 2016, 61, 125-137.	4.5	123
5	Crystal Structure of the Herpesvirus Nuclear Egress Complex Provides Insights into Inner Nuclear Membrane Remodeling. Cell Reports, 2015, 13, 2645-2652.	2.9	80
6	The crystal structure of human dopamine \hat{l}^2 -hydroxylase at 2.9 \tilde{A} resolution. Science Advances, 2016, 2, e1500980.	4.7	80
7	Unexpected structure for the N-terminal domain of hepatitis C virus envelope glycoprotein E1. Nature Communications, 2014, 5, 4874.	5.8	72
8	Unexpected features and mechanism of heterodimer formation of a herpesvirus nuclear egress complex. EMBO Journal, 2015, 34, 2937-2952.	3.5	69
9	Structural basis for antibacterial peptide selfâ€immunity by the bacterial ABC transporter McjD. EMBO Journal, 2017, 36, 3062-3079.	3.5	64
10	Molecular Architecture and Ligand Recognition Determinants for T4 RNA Ligase. Journal of Biological Chemistry, 2006, 281, 1573-1579.	1.6	61
11	Structure of the leukemia oncogene LMO2: implications for the assembly of a hematopoietic transcription factor complex. Blood, 2011, 117, 2146-2156.	0.6	59
12	Multiple liquid crystalline geometries of highly compacted nucleic acid in a dsRNA virus. Nature, 2019, 570, 252-256.	13.7	59
13	Structural Basis for LMO2-Driven Recruitment of the SCL:E47bHLH Heterodimer to Hematopoietic-Specific Transcriptional Targets. Cell Reports, 2013, 4, 135-147.	2.9	56
14	The morphogen Sonic hedgehog inhibits its receptor Patched by a pincer grasp mechanism. Nature Chemical Biology, 2019, 15, 975-982.	3.9	52
15	The conserved protein Seb1 drives transcription termination by binding RNA polymerase II and nascent RNA. Nature Communications, 2017, 8, 14861.	5.8	48
16	Plate Tectonics of Virus Shell Assembly and Reorganization in Phage $\hat{l} \mid 8$, a Distant Relative of Mammalian Reoviruses. Structure, 2013, 21, 1384-1395.	1.6	45
17	Anomalous X-ray diffraction studies of ion transport in K+ channels. Nature Communications, 2018, 9, 4540.	5.8	42
18	Structural basis for non-competitive product inhibition in human thymidine phosphorylase: implications for drug design. Biochemical Journal, 2006, 399, 199-204.	1.7	38

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19	The structure of a prokaryotic viral envelope protein expands the landscape of membrane fusion proteins. Nature Communications, 2019, 10, 846.	5.8	37
20	Double-stranded RNA virus outer shell assembly by bona fide domain-swapping. Nature Communications, 2017, 8, 14814.	5.8	35
21	Pushing the limits of sulfur SAD phasing: <i>de novo</i> structure solution of the N-terminal domain of the ectodomain of HCV E1. Acta Crystallographica Section D: Biological Crystallography, 2014, 70, 2197-2203.	2.5	33
22	Structure of vaccinia virus thymidine kinase in complex with dTTP: insights for drug design. BMC Structural Biology, 2006, 6, 22.	2.3	31
23	Structure and assembly of the S-layer in C. difficile. Nature Communications, 2022, 13, 970.	5.8	30
24	Crystal Structure of CC3 (TIP30). Journal of Biological Chemistry, 2005, 280, 18229-18236.	1.6	27
25	Tracking in atomic detail the functional specializations in viral RecA helicases that occur during evolution. Nucleic Acids Research, 2013, 41, 9396-9410.	6.5	23
26	Structure of Staphylococcus aureus guanylate monophosphate kinase. Acta Crystallographica Section F: Structural Biology Communications, 2006, 62, 949-953.	0.7	21
27	Structure of the DNAâ€bound Tâ€box domain of human TBX1, a transcription factor associated with the DiGeorge syndrome. Proteins: Structure, Function and Bioinformatics, 2012, 80, 655-660.	1.5	21
28	Mutations Distal to the Substrate Site Can Affect Varicella Zoster Virus Thymidine Kinase Activity: Implications for Drug Design. Molecular Pharmacology, 2006, 69, 1891-1896.	1.0	16
29	Hedgehog-Interacting Protein is a multimodal antagonist of Hedgehog signalling. Nature Communications, 2021, 12, 7171.	5.8	16
30	Functional metagenomic screening identifies an unexpected \hat{l}^2 -glucuronidase. Nature Chemical Biology, 2022, 18, 1096-1103.	3.9	16
31	Native de novo structural determinations of non-canonical nucleic acid motifs by X-ray crystallography at long wavelengths. Nucleic Acids Research, 2020, 48, 9886-9898.	6.5	13
32	Atypical Porcine Pestiviruses: Relationships and Conserved Structural Features. Viruses, 2021, 13, 760.	1.5	5
33	Identifying dynamic, partially occupied residues using anomalous scattering. Acta Crystallographica Section D: Structural Biology, 2019, 75, 1084-1095.	1.1	5
34	Purification, crystallization and preliminary X-ray analysis of a fusion of the LIM domains of LMO2 and the LID domain of Ldb1. Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 1466-1469.	0.7	4
35	Experimental phasing with vanadium and application to nucleotide-binding membrane proteins. IUCrJ, 2020, 7, 1092-1101.	1.0	3
36	Phosphorus and sulfur SAD phasing of the nucleic acid-bound DNA-binding domain of interferon regulatory factor 4. Acta Crystallographica Section F, Structural Biology Communications, 2021, 77, 202-207.	0.4	2

#	Article	IF	CITATIONS
37	The design and development of drugs acting against the smallpox virus. Expert Opinion on Drug Discovery, 2007, 2, 1263-1272.	2.5	1