

Arnaud BaslÃ©

List of Publications by Year in descending order

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Version: 2024-02-01

29
papers

1,478
citations

516710

16
h-index

477307

29
g-index

30
all docs

30
docs citations

30
times ranked

1996
citing authors

#	ARTICLE	IF	CITATIONS
1	Glycan complexity dictates microbial resource allocation in the large intestine. <i>Nature Communications</i> , 2015, 6, 7481.	12.8	328
2	Structural basis for nutrient acquisition by dominant members of the human gut microbiota. <i>Nature</i> , 2017, 541, 407-411.	27.8	188
3	Structural basis for maintenance of bacterial outer membrane lipid asymmetry. <i>Nature Microbiology</i> , 2017, 2, 1616-1623.	13.3	118
4	Complex N-glycan breakdown by gut <i>Bacteroides</i> involves an extensive enzymatic apparatus encoded by multiple co-regulated genetic loci. <i>Nature Microbiology</i> , 2019, 4, 1571-1581.	13.3	116
5	A four-helix bundle stores copper for methane oxidation. <i>Nature</i> , 2015, 525, 140-143.	27.8	83
6	Prominent members of the human gut microbiota express endo-acting O-glycanases to initiate mucin breakdown. <i>Nature Communications</i> , 2020, 11, 4017.	12.8	81
7	Small-Molecule Transport by CarO, an Abundant Eight-Stranded β^2 -Barrel Outer Membrane Protein from <i>Acinetobacter baumannii</i> . <i>Journal of Molecular Biology</i> , 2015, 427, 2329-2339.	4.2	54
8	Bacterial cytosolic proteins with a high capacity for Cu(I) that protect against copper toxicity. <i>Scientific Reports</i> , 2016, 6, 39065.	3.3	52
9	Structural basis for Mep2 ammonium transceptor activation by phosphorylation. <i>Nature Communications</i> , 2016, 7, 11337.	12.8	52
10	Structural Insights into Outer Membrane Permeability of <i>Acinetobacter baumannii</i> . <i>Structure</i> , 2016, 24, 221-231.	3.3	49
11	Structural and functional insights into oligopeptide acquisition by the RagAB transporter from <i>Porphyromonas gingivalis</i> . <i>Nature Microbiology</i> , 2020, 5, 1016-1025.	13.3	46
12	Insights into SusCD-mediated glycan import by a prominent gut symbiont. <i>Nature Communications</i> , 2021, 12, 44.	12.8	42
13	Structural basis for chitin acquisition by marine <i>Vibrio</i> species. <i>Nature Communications</i> , 2018, 9, 220.	12.8	37
14	Structure and assembly of the S-layer in <i>C. difficile</i> . <i>Nature Communications</i> , 2022, 13, 970.	12.8	30
15	Unusual Constriction Zones in the Major Porins OmpU and OmpT from <i>Vibrio cholerae</i> . <i>Structure</i> , 2018, 26, 708-721.e4.	3.3	22
16	An evolutionary path to altered cofactor specificity in a metalloenzyme. <i>Nature Communications</i> , 2020, 11, 2738.	12.8	22
17	Pore dynamics and asymmetric cargo loading in an encapsulin nanocompartment. <i>Science Advances</i> , 2022, 8, eabj4461.	10.3	22
18	Crystal structure of NucB, a biofilm-degrading endonuclease. <i>Nucleic Acids Research</i> , 2018, 46, 473-484.	14.5	21

#	ARTICLE	IF	CITATIONS
19	Structural Basis for Silicic Acid Uptake by Higher Plants. <i>Journal of Molecular Biology</i> , 2021, 433, 167226.	4.2	18
20	Visualizing Biological Copper Storage: The Importance of Thiolateâ€Coordinated Tetranuclear Clusters. <i>Angewandte Chemie - International Edition</i> , 2017, 56, 8697-8700.	13.8	16
21	Insight into Metal Removal from Peptides that Sequester Copper for Methane Oxidation. <i>Chemistry - A European Journal</i> , 2018, 24, 4515-4518.	3.3	16
22	Acquisition of ionic copper by the bacterial outer membrane protein OprC through a novel binding site. <i>PLoS Biology</i> , 2021, 19, e3001446.	5.6	14
23	Dissecting the structural and functional roles of a putative metal entry site in encapsulated ferritins. <i>Journal of Biological Chemistry</i> , 2020, 295, 15511-15526.	3.4	13
24	Uptake of monoaromatic hydrocarbons during biodegradation by FadL channel-mediated lateral diffusion. <i>Nature Communications</i> , 2020, 11, 6331.	12.8	10
25	Crystal structure of the <i>Acinetobacter baumannii</i> outer membrane protein Omp33. <i>Acta Crystallographica Section D: Structural Biology</i> , 2018, 74, 852-860.	2.3	8
26	The predictive power of data-processing statistics. <i>IUCr</i> , 2020, 7, 342-354.	2.2	7
27	A comprehensive structural analysis of the ATPase domain of human DNA topoisomerase II beta bound to AMPPNP, ADP, and the bisdioxopiperazine, ICRF193. <i>Structure</i> , 2022, 30, 1129-1145.e3.	3.3	6
28	Visualizing Biological Copper Storage: The Importance of Thiolateâ€Coordinated Tetranuclear Clusters. <i>Angewandte Chemie</i> , 2017, 129, 8823-8826.	2.0	3
29	A practical overview of molecular replacement: <i>Clostridioides difficile</i> PilA1, a difficult case study. <i>Acta Crystallographica Section D: Structural Biology</i> , 2020, 76, 261-271.	2.3	3