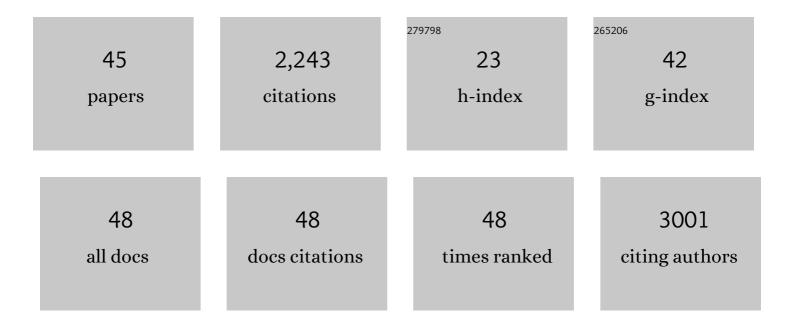
Roman Brunecky

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The Structural Origin of Chiroptical Properties in Perovskite Nanocrystals with Chiral Organic Ligands. Advanced Functional Materials, 2022, 32, .	14.9	43
2	A Swollenin From Talaromyces leycettanus JCM12802 Enhances Cellulase Hydrolysis Toward Various Substrates. Frontiers in Microbiology, 2021, 12, 658096.	3.5	7
3	Substitution of distal and active site residues reduces product inhibition of E1 from <i>Acidothermus Cellulolyticus</i> . Protein Engineering, Design and Selection, 2021, 34, .	2.1	2
4	Synthetic fungal multifunctional cellulases for enhanced biomass conversion. Green Chemistry, 2020, 22, 478-489.	9.0	31
5	Highly Distorted Chiral Two-Dimensional Tin Iodide Perovskites for Spin Polarized Charge Transport. Journal of the American Chemical Society, 2020, 142, 13030-13040.	13.7	198
6	Strategies to Achieve High Circularly Polarized Luminescence from Colloidal Organic–Inorganic Hybrid Perovskite Nanocrystals. ACS Nano, 2020, 14, 8816-8825.	14.6	94
7	Glycosylation Is Vital for Industrial Performance of Hyperactive Cellulases. ACS Sustainable Chemistry and Engineering, 2019, 7, 4792-4800.	6.7	19
8	Spin-dependent charge transport through 2D chiral hybrid lead-iodide perovskites. Science Advances, 2019, 5, eaay0571.	10.3	275
9	Handling gene and protein names in the age of bioinformatics: the special challenge of secreted multimodular bacterial enzymes such as the cbhA/cbh9A gene of Clostridium thermocellum. World Journal of Microbiology and Biotechnology, 2018, 34, 42.	3.6	2
10	High activity CAZyme cassette for improving biomass degradation in thermophiles. Biotechnology for Biofuels, 2018, 11, 22.	6.2	35
11	The Multi Domain Caldicellulosiruptor bescii CelA Cellulase Excels at the Hydrolysis of Crystalline Cellulose. Scientific Reports, 2017, 7, 9622.	3.3	43
12	Biomass Conversion. , 2017, , 285-419.		7
13	Towards an Understanding of Enhanced Biomass Digestibility by In Planta Expression of a Family 5 Glycoside Hydrolase. Scientific Reports, 2017, 7, 4389.	3.3	9
14	Natural diversity of glycoside hydrolase family 48 exoglucanases: insights from structure. Biotechnology for Biofuels, 2017, 10, 274.	6.2	7
15	Undefined cellulase formulations hinder scientific reproducibility. Biotechnology for Biofuels, 2017, 10, 283.	6.2	7
16	Strategies to reduce endâ€product inhibition in family 48 glycoside hydrolases. Proteins: Structure, Function and Bioinformatics, 2016, 84, 295-304.	2.6	10
17	Crystal structure and biochemical characterization of Chlamydomonas FDX2 reveal two residues that, when mutated, partially confer FDX2 the redox potential and catalytic properties of FDX1. Photosynthesis Research, 2016, 128, 45-57.	2.9	22
18	The catalytic mechanism and unique low pH optimum of <i>Caldicellulosiruptor bescii</i> family 3 pectate lyase. Acta Crystallographica Section D: Biological Crystallography, 2015, 71, 1946-1954.	2.5	14

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19	Identifying the ionically bound cell wall and intracellular glycoside hydrolases in late growth stage Arabidopsis stems: implications for the genetic engineering of bioenergy crops. Frontiers in Plant Science, 2015, 6, 315.	3.6	14
20	New Insights into Microbial Strategies for Biomass Conversion. , 2015, , 111-127.		4
21	Expression of the Acidothermus cellulolyticus E1 endoglucanase in Caldicellulosiruptor bescii enhances its ability to deconstruct crystalline cellulose. Biotechnology for Biofuels, 2015, 8, 113.	6.2	31
22	Investigation of the role of lignin in biphasic xylan hydrolysis during dilute acid and organosolv pretreatment of corn stover. Green Chemistry, 2015, 17, 1546-1558.	9.0	20
23	Feedstock Engineering and Biomass Pretreatments. , 2015, , 3-12.		1
24	High temperature pre-digestion of corn stover biomass for improved product yields. Biotechnology for Biofuels, 2014, 7, 170.	6.2	11
25	Cel48A from <i>Thermobifida fusca</i> : Structure and site directed mutagenesis of key residues. Biotechnology and Bioengineering, 2014, 111, 664-673.	3.3	35
26	Charge engineering of cellulases improves ionic liquid tolerance and reduces lignin inhibition. Biotechnology and Bioengineering, 2014, 111, 1541-1549.	3.3	91
27	Response to Comment on "Revealing Nature's Cellulase Diversity: The Digestion Mechanism of <i>Caldicellulosiruptor bescii</i> CelAâ€: Science, 2014, 344, 578-578.	12.6	1
28	Revealing Nature's Cellulase Diversity: The Digestion Mechanism of <i>Caldicellulosiruptor bescii</i> CelA. Science, 2013, 342, 1513-1516.	12.6	253
29	Improving activity of minicellulosomes by integration of intra- and intermolecular synergies. Biotechnology for Biofuels, 2013, 6, 126.	6.2	37
30	Impact of alg3 gene deletion on growth, development, pigment production, protein secretion, and functions of recombinant Trichoderma reesei cellobiohydrolases in Aspergillus niger. Fungal Genetics and Biology, 2013, 61, 120-132.	2.1	25
31	The structure and mode of action ofCaldicellulosiruptor besciifamily 3 pectate lyase in biomass deconstruction. Acta Crystallographica Section D: Biological Crystallography, 2013, 69, 534-539.	2.5	7
32	Sequence, Structure, and Evolution of Cellulases in Glycoside Hydrolase Family 48. Journal of Biological Chemistry, 2012, 287, 41068-41077.	3.4	32
33	Structure and function of theClostridium thermocellumcellobiohydrolase A X1-module repeat: enhancement through stabilization of the CbhA complex. Acta Crystallographica Section D: Biological Crystallography, 2012, 68, 292-299.	2.5	15
34	Analysis of Transgenic Glycoside Hydrolases Expressed in Plants: T. reesei CBH I and A. cellulolyticus El. Methods in Molecular Biology, 2012, 908, 197-211.	0.9	3
35	Cellulase Linkers Are Optimized Based on Domain Type and Function: Insights from Sequence Analysis, Biophysical Measurements, and Molecular Simulation. PLoS ONE, 2012, 7, e48615.	2.5	88
36	Bioprospecting metagenomics of decaying wood: mining for new glycoside hydrolases. Biotechnology for Biofuels, 2011, 4, 23.	6.2	40

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#	Article	IF	CITATIONS
37	In planta expression of A. cellulolyticus Cel5A endocellulase reduces cell wall recalcitrance in tobacco and maize. Biotechnology for Biofuels, 2011, 4, 1.	6.2	217
38	Structure of a fibronectin type III-like module fromClostridium thermocellum. Acta Crystallographica Section F: Structural Biology Communications, 2010, 66, 878-880.	0.7	12
39	The Unique Binding Mode of Cellulosomal CBM4 from Clostridium thermocellum Cellobiohydrolase A. Journal of Molecular Biology, 2010, 402, 374-387.	4.2	28
40	ORIGINAL RESEARCH: Lignocellulose recalcitrance screening by integrated high-throughput hydrothermal pretreatment and enzymatic saccharification. Industrial Biotechnology, 2010, 6, 104-111.	0.8	80
41	Redistribution of xylan in maize cell walls during dilute acid pretreatment. Biotechnology and Bioengineering, 2009, 102, 1537-1543.	3.3	53
42	Probing the role of N-linked glycans in the stability and activity of fungal cellobiohydrolases by mutational analysis. Cellulose, 2009, 16, 699-709.	4.9	79
43	High-Throughput Screening Techniques for Biomass Conversion. Bioenergy Research, 2009, 2, 179-192.	3.9	82
44	Heterologous expression of glycosyl hydrolases in planta: a new departure for biofuels. Trends in Biotechnology, 2008, 26, 413-424.	9.3	115
45	Investigation of the Binding Geometry of a Peripheral Membrane Proteinâ€. Biochemistry, 2005, 44, 16064-16071.	2.5	34