

# Robin W Palfreyman

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2916444/publications.pdf>

Version: 2024-02-01

25  
papers

1,902  
citations

516710

16  
h-index

610901

24  
g-index

26  
all docs

26  
docs citations

26  
times ranked

3247  
citing authors

#	ARTICLE	IF	CITATIONS
1	Multi-omic characterisation of <i>Streptomyces hygrosopicus</i> NRRL 30439: detailed assessment of its secondary metabolic potential. <i>Molecular Omics</i> , 2022, 18, 226-236.	2.8	5
2	Role of the substrate on Ni inhibition in biological sulfate reduction. <i>Journal of Environmental Management</i> , 2022, 316, 115216.	7.8	0
3	Network Analyses Predict Small RNAs That Might Modulate Gene Expression in the Testis and Epididymis of <i>Bos indicus</i> Bulls. <i>Frontiers in Genetics</i> , 2021, 12, 610116.	2.3	7
4	A Pan-Genome Guided Metabolic Network Reconstruction of Five <i>Propionibacterium</i> Species Reveals Extensive Metabolic Diversity. <i>Genes</i> , 2020, 11, 1115.	2.4	18
5	A snapshot of microbial diversity and function in an undisturbed sugarcane bagasse pile. <i>BMC Biotechnology</i> , 2020, 20, 12.	3.3	12
6	A TetR-Family Protein (CAETHG_0459) Activates Transcription From a New Promoter Motif Associated With Essential Genes for Autotrophic Growth in Acetogens. <i>Frontiers in Microbiology</i> , 2019, 10, 2549.	3.5	12
7	Systems-level engineering and characterisation of <i>Clostridium autoethanogenum</i> through heterologous production of poly-3-hydroxybutyrate (PHB). <i>Metabolic Engineering</i> , 2019, 53, 14-23.	7.0	57
8	From reconstruction to C4 metabolic engineering: A case study for overproduction of polyhydroxybutyrate in bioenergy grasses. <i>Plant Science</i> , 2018, 273, 50-60.	3.6	7
9	RNA-Seq Highlights High Clonal Variation in Monoclonal Antibody Producing CHO Cells. <i>Biotechnology Journal</i> , 2018, 13, e1700231.	3.5	28
10	Effect of Plasmid Design and Type of Integration Event on Recombinant Protein Expression in <i>Pichia pastoris</i> . <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	54
11	Maintenance of ATP Homeostasis Triggers Metabolic Shifts in Gas-Fermenting Acetogens. <i>Cell Systems</i> , 2017, 4, 505-515.e5.	6.2	128
12	Improved production of propionic acid using genome shuffling. <i>Biotechnology Journal</i> , 2017, 12, 1600120.	3.5	23
13	Metabolic Reconstruction of <i>Setaria italica</i> : A Systems Biology Approach for Integrating Tissue-Specific Omics and Pathway Analysis of Bioenergy Grasses. <i>Frontiers in Plant Science</i> , 2016, 7, 1138.	3.6	24
14	Genomic characterization of the uncultured Bacteroidales family S24-7 inhabiting the guts of homeothermic animals. <i>Microbiome</i> , 2016, 4, 36.	11.1	533
15	Systems biology and metabolic modelling unveils limitations to polyhydroxybutyrate accumulation in sugarcane leaves; lessons for <i>SCP</i> engineering. <i>Plant Biotechnology Journal</i> , 2016, 14, 567-580.	8.3	17
16	Low carbon fuels and commodity chemicals from waste gases – systematic approach to understand energy metabolism in a model acetogen. <i>Green Chemistry</i> , 2016, 18, 3020-3028.	9.0	143
17	Evolutionary Engineering Improves Tolerance for Replacement Jet Fuels in <i>Saccharomyces cerevisiae</i> . <i>Applied and Environmental Microbiology</i> , 2015, 81, 3316-3325.	3.1	44
18	Plant Genome-Scale Modeling and Implementation. <i>Methods in Molecular Biology</i> , 2014, 1090, 317-332.	0.9	8

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19	Saccharopolyspora erythraea genome is organised in high-order transcriptional regions mediated by targeted degradation at the metabolic switch. BMC Genomics, 2013, 14, 15.	2.8	33
20	Re-annotation of the Saccharopolyspora erythraea genome using a systems biology approach. BMC Genomics, 2013, 14, 699.	2.8	21
21	Transcriptome Sequencing of and Microarray Development for a Helicoverpa zea Cell Line to Investigate In Vitro Insect Cell-Baculovirus Interactions. PLoS ONE, 2012, 7, e36324.	2.5	28
22	Deep sequencing-based transcriptome analysis of Plutella xylostella larvae parasitized by Diadegma semiclausum. BMC Genomics, 2011, 12, 446.	2.8	82
23	AlgaGEM – a genome-scale metabolic reconstruction of algae based on the Chlamydomonas reinhardtii genome. BMC Genomics, 2011, 12, S5.	2.8	109
24	AraGEM, a Genome-Scale Reconstruction of the Primary Metabolic Network in Arabidopsis. Plant Physiology, 2010, 152, 579-589.	4.8	319
25	C4GEM, a Genome-Scale Metabolic Model to Study C4 Plant Metabolism. Plant Physiology, 2010, 154, 1871-1885.	4.8	190