

# Piotr Grabowski

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/2155879/publications.pdf>

Version: 2024-02-01

8  
papers

276  
citations

1684188

5  
h-index

2053705

5  
g-index

11  
all docs

11  
docs citations

11  
times ranked

626  
citing authors

#	ARTICLE	IF	CITATIONS
1	Co-regulation map of the human proteome enables identification of protein functions. <i>Nature Biotechnology</i> , 2019, 37, 1361-1371.	17.5	106
2	Pervasive coexpression of spatially proximal genes is buffered at the protein level. <i>Molecular Systems Biology</i> , 2017, 13, 937.	7.2	90
3	Proteome Analysis of Human Neutrophil Granulocytes From Patients With Monogenic Disease Using Data-independent Acquisition. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 760-772.	3.8	52
4	A Primer on Data Analytics in Functional Genomics: How to Move from Data to Insight?. <i>Trends in Biochemical Sciences</i> , 2019, 44, 21-32.	7.5	16
5	Epigenetic Variability Confounds Transcriptome but Not Proteome Profiling for Coexpression-based Gene Function Prediction. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 2082-2090.	3.8	9
6	Defects in Signal Recognition Particle (SRP) Components Reveal an Essential and Non-Redundant Role for Granule Biogenesis and Differentiation of Neutrophil Granulocytes. <i>Blood</i> , 2019, 134, 216-216.	1.4	1
7	Next Generation Proteomics of Human Neutrophil Granulocytes in Monogenic Disease. <i>Blood</i> , 2018, 132, 15-15.	1.4	0
8	Machine Learning Unveils Proteotypic Mimicry in Genetically Defined SCN Variants. <i>Blood</i> , 2019, 134, 3580-3580.	1.4	0