

Marco L Hennrich

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8547636/publications.pdf>

Version: 2024-02-01

9
papers

447
citations

1477746

6
h-index

1588620

8
g-index

10
all docs

10
docs citations

10
times ranked

1153
citing authors

#	ARTICLE	IF	CITATIONS
1	Impact of spliceosome mutations on RNA splicing in myelodysplasia: dysregulated genes/pathways and clinical associations. <i>Blood</i> , 2018, 132, 1225-1240.	0.6	168
2	Capturing protein communities by structural proteomics in a thermophilic eukaryote. <i>Molecular Systems Biology</i> , 2017, 13, 936.	3.2	108
3	Cell-specific proteome analyses of human bone marrow reveal molecular features of age-dependent functional decline. <i>Nature Communications</i> , 2018, 9, 4004.	5.8	71
4	Phosphatase of regenerating liver (PRL)-3 disrupts epithelial architecture by altering the post-mitotic midbody position. <i>Journal of Cell Science</i> , 2016, 129, 4130-4142.	1.2	33
5	Cell metabolism regulates integrin mechanosensing via an SLC3A2-dependent sphingolipid biosynthesis pathway. <i>Nature Communications</i> , 2018, 9, 4862.	5.8	28
6	Dynamic Buffering of Extracellular Chemokine by a Dedicated Scavenger Pathway Enables Robust Adaptation during Directed Tissue Migration. <i>Developmental Cell</i> , 2020, 52, 492-508.e10.	3.1	25
7	CLICK-enabled analogues reveal pregnenolone interactomes in cancer and immune cells. <i>IScience</i> , 2021, 24, 102485.	1.9	6
8	Coupling proteomics and metabolomics for the unsupervised identification of protein-metabolite interactions in <i>Chaetomium thermophilum</i> . <i>PLoS ONE</i> , 2021, 16, e0254429.	1.1	5
9	Proteome Analyses and Single-Cell RNA Sequencing Reveal Age-Dependent Re-Wiring of Central Carbon Metabolism in Myeloid-Biased Subsets of Human Hematopoietic Stem Cells. <i>Blood</i> , 2018, 132, 873-873.	0.6	0