

# Jonathan Martinez-Fabregas

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/387193/publications.pdf>

Version: 2024-02-01

15  
papers

448  
citations

933447

10  
h-index

1281871

11  
g-index

18  
all docs

18  
docs citations

18  
times ranked

719  
citing authors

#	ARTICLE	IF	CITATIONS
1	PD-1 Inhibitory Receptor Downregulates Asparaginyl Endopeptidase and Maintains Foxp3 Transcription Factor Stability in Induced Regulatory T Cells. <i>Immunity</i> , 2018, 49, 247-263.e7.	14.3	104
2	Structural and Functional Analysis of Novel Human Cytochrome c Targets in Apoptosis. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 1439-1456.	3.8	74
3	New Arabidopsis thaliana Cytochrome c Partners: A Look Into the Elusive Role of Cytochrome c in Programmed Cell Death in Plants. <i>Molecular and Cellular Proteomics</i> , 2013, 12, 3666-3676.	3.8	58
4	Lysosomal protease deficiency or substrate overload induces an oxidative-stress mediated STAT3-dependent pathway of lysosomal homeostasis. <i>Nature Communications</i> , 2018, 9, 5343.	12.8	52
5	A common signalosome for programmed cell death in humans and plants. <i>Cell Death and Disease</i> , 2014, 5, e1314-e1314.	6.3	41
6	Kinetics of cytokine receptor trafficking determine signaling and functional selectivity. <i>ELife</i> , 2019, 8, .	6.0	34
7	CDK8 Fine-Tunes IL-6 Transcriptional Activities by Limiting STAT3 Resident Time at the Gene Loci. <i>Cell Reports</i> , 2020, 33, 108545.	6.4	26
8	Competitive binding of STATs to receptor phospho-Tyr motifs accounts for altered cytokine responses. <i>ELife</i> , 2021, 10, .	6.0	21
9	Mapping Determinants of Cytokine Signaling via Protein Engineering. <i>Frontiers in Immunology</i> , 2018, 9, 2143.	4.8	20
10	Proteomic tools for the analysis of transient interactions between metalloproteins. <i>FEBS Journal</i> , 2011, 278, 1401-1410.	4.7	12
11	Lysosomes: multifunctional compartments ruled by a complex regulatory network. <i>FEBS Open Bio</i> , 2022, 12, 758-774.	2.3	3
12	Structural and functional changes induced by tyrosine nitration in cytochrome c, a bi-functional protein. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2010, 1797, 70.	1.0	0
13	Identifying cytokine signaling signatures in primary human Th-1 cells by phospho-proteomics analysis. <i>STAR Protocols</i> , 2021, 2, 100417.	1.2	0
14	A Common Cytochrome c -centred Signalosome for Programmed Cell Death in Humans and Plants. <i>FASEB Journal</i> , 2015, 29, 569.13.	0.5	0
15	Cytochrome c -Based Signalosome. , 2015, , 275-298.		0