

# Pierre Sabatier

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3704687/publications.pdf>

Version: 2024-02-01

13  
papers

427  
citations

1170033

9  
h-index

1255698

13  
g-index

20  
all docs

20  
docs citations

20  
times ranked

531  
citing authors

#	ARTICLE	IF	CITATIONS
1	A "spindle and thread" mechanism unblocks p53 translation by modulating N-terminal disorder. <i>Structure</i> , 2022, 30, 733-742.e7.	1.6	5
2	Comprehensive Target Screening and Cellular Profiling of the Cancer-Active Compound b-AP15 Indicate Abrogation of Protein Homeostasis and Organelle Dysfunction as the Primary Mechanism of Action. <i>Frontiers in Oncology</i> , 2022, 12, 852980.	1.3	2
3	Ion-Based Proteome-Integrated Solubility Alteration Assays for Systemwide Profiling of Protein-Molecule Interactions. <i>Analytical Chemistry</i> , 2022, 94, 7066-7074.	3.2	12
4	System-wide identification and prioritization of enzyme substrates by thermal analysis. <i>Nature Communications</i> , 2021, 12, 1296.	5.8	44
5	Polymorphic estrogen receptor binding site causes Cd2-dependent sex bias in the susceptibility to autoimmune diseases. <i>Nature Communications</i> , 2021, 12, 5565.	5.8	12
6	An integrative proteomics method identifies a regulator of translation during stem cell maintenance and differentiation. <i>Nature Communications</i> , 2021, 12, 6558.	5.8	16
7	Comprehensive chemical proteomics analyses reveal that the new TRi-1 and TRi-2 compounds are more specific thioredoxin reductase 1 inhibitors than auranofin. <i>Redox Biology</i> , 2021, 48, 102184.	3.9	18
8	Vitamin D3 receptor polymorphisms regulate T cells and T cell-dependent inflammatory diseases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 24986-24997.	3.3	14
9	Comprehensive chemical proteomics for target deconvolution of the redox active drug auranofin. <i>Redox Biology</i> , 2020, 32, 101491.	3.9	58
10	Proteome Integral Solubility Alteration: A High-Throughput Proteomics Assay for Target Deconvolution. <i>Journal of Proteome Research</i> , 2019, 18, 4027-4037.	1.8	148
11	ProTargetMiner as a proteome signature library of anticancer molecules for functional discovery. <i>Nature Communications</i> , 2019, 10, 5715.	5.8	47
12	Comparative Proteomics of Dying and Surviving Cancer Cells Improves the Identification of Drug Targets and Sheds Light on Cell Life/Death Decisions. <i>Molecular and Cellular Proteomics</i> , 2018, 17, 1144-1155.	2.5	25
13	Dynamic Proteomics Reveals High Plasticity of Cellular Proteome: Growth-Related and Drug-Induced Changes in Cancer Cells are Comparable. <i>Proteomics</i> , 2018, 18, e1800118.	1.3	14