

Lingyan Ping

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3651912/publications.pdf>

Version: 2024-02-01

13
papers

1,376
citations

1163117

8
h-index

1281871

11
g-index

17
all docs

17
docs citations

17
times ranked

1751
citing authors

#	ARTICLE	IF	CITATIONS
1	Large-scale proteomic analysis of Alzheimer's disease brain and cerebrospinal fluid reveals early changes in energy metabolism associated with microglia and astrocyte activation. <i>Nature Medicine</i> , 2020, 26, 769-780.	30.7	547
2	Large-scale deep multi-layer analysis of Alzheimer's disease brain reveals strong proteomic disease-related changes not observed at the RNA level. <i>Nature Neuroscience</i> , 2022, 25, 213-225.	14.8	202
3	Integrated proteomics reveals brain-based cerebrospinal fluid biomarkers in asymptomatic and symptomatic Alzheimer's disease. <i>Science Advances</i> , 2020, 6, .	10.3	186
4	Global quantitative analysis of the human brain proteome in Alzheimer's and Parkinson's Disease. <i>Scientific Data</i> , 2018, 5, 180036.	5.3	179
5	Global quantitative analysis of the human brain proteome and phosphoproteome in Alzheimer's disease. <i>Scientific Data</i> , 2020, 7, 315.	5.3	74
6	Targeted mass spectrometry to quantify brain-derived cerebrospinal fluid biomarkers in Alzheimer's disease. <i>Clinical Proteomics</i> , 2020, 17, 19.	2.1	53
7	Quantitative Analysis of the Brain Ubiquitylome in Alzheimer's Disease. <i>Proteomics</i> , 2018, 18, e1800108.	2.2	50
8	Extracellular signal-regulated kinase regulates microglial immune responses in Alzheimer's disease. <i>Journal of Neuroscience Research</i> , 2021, 99, 1704-1721.	2.9	43
9	Signatures of glial activity can be detected in the CSF proteome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	12
10	Phosphorylation regulates arginine-rich RNA-binding protein solubility and oligomerization. <i>Journal of Biological Chemistry</i> , 2021, 297, 101306.	3.4	9
11	Large-scale deep multi-layer analysis of Alzheimer's disease brain reveals strong proteomic disease-related changes not observed at the RNA level. <i>Alzheimer's and Dementia</i> , 2021, 17, e055041.	0.8	1
12	A consensus proteomic analysis of Alzheimer's disease brain and cerebrospinal fluid reveals early changes in energy metabolism associated with microglia and astrocyte activation. <i>Alzheimer's and Dementia</i> , 2020, 16, e039504.	0.8	0
13	Proteomics identifies CSF biomarker panels reflective of pathological networks in the Alzheimer's disease brain. <i>Alzheimer's and Dementia</i> , 2020, 16, e042227.	0.8	0