

Qifeng Song

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10877891/publications.pdf>

Version: 2024-02-01

13
papers

820
citations

687363

13
h-index

1125743

13
g-index

17
all docs

17
docs citations

17
times ranked

1855
citing authors

#	ARTICLE	IF	CITATIONS
1	DNA methylation presents distinct binding sites for human transcription factors. <i>ELife</i> , 2013, 2, e00726.	6.0	292
2	Lin28A Binds Active Promoters and Recruits Tet1 to Regulate Gene Expression. <i>Molecular Cell</i> , 2016, 61, 153-160.	9.7	74
3	Novel Autoimmune Hepatitis-Specific Autoantigens Identified Using Protein Microarray Technology. <i>Journal of Proteome Research</i> , 2010, 9, 30-39.	3.7	71
4	Sensitive Affimer and Antibody Based Impedimetric Label-Free Assays for C-Reactive Protein. <i>Analytical Chemistry</i> , 2012, 84, 6553-6560.	6.5	68
5	Label-Free Sub-picomolar Protein Detection with Field-Effect Transistors. <i>Analytical Chemistry</i> , 2010, 82, 3531-3536.	6.5	61
6	Functional Characterization of Schizophrenia-Associated Variation in CACNA1C. <i>PLoS ONE</i> , 2016, 11, e0157086.	2.5	61
7	Structure-function studies of an engineered scaffold protein derived from Stefin A. II: Development and applications of the SQT variant. <i>Protein Engineering, Design and Selection</i> , 2011, 24, 751-763.	2.1	43
8	Methylated cis-regulatory elements mediate KLF4-dependent gene transactivation and cell migration. <i>ELife</i> , 2017, 6, .	6.0	39
9	Structure-function studies of an engineered scaffold protein derived from stefin A. I: Development of the SQM variant. <i>Protein Engineering, Design and Selection</i> , 2010, 23, 403-413.	2.1	31
10	A protein chip approach for high-throughput antigen identification and characterization. <i>Proteomics</i> , 2007, 7, 2151-2161.	2.2	24
11	An all-to-all approach to the identification of sequence-specific readers for epigenetic DNA modifications on cytosine. <i>Nature Communications</i> , 2021, 12, 795.	12.8	22
12	Proteome-wide Tyrosine Phosphorylation Analysis Reveals Dysregulated Signaling Pathways in Ovarian Tumors. <i>Molecular and Cellular Proteomics</i> , 2019, 18, 448-460.	3.8	19
13	Peptide Aptamer Microarrays: Bridging the bio-sensor detector interface. <i>Faraday Discussions</i> , 2011, 149, 79-92.	3.2	14