## Qifeng Song

List of Publications by Year in descending order

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

Version: 2024-02-01

1125743
13
g-index
1055
1855
citing authors

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