

# Jun Seop Jeong

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12023684/publications.pdf>

Version: 2024-02-01

14  
papers

2,510  
citations

840776

11  
h-index

1058476

14  
g-index

15  
all docs

15  
docs citations

15  
times ranked

4467  
citing authors

#	ARTICLE	IF	CITATIONS
1	The genome sequence of the rice blast fungus <i>Magnaporthe grisea</i> . <i>Nature</i> , 2005, 434, 980-986.	27.8	1,447
2	A nuclease that mediates cell death induced by DNA damage and poly(ADP-ribose) polymerase-1. <i>Science</i> , 2016, 354, .	12.6	266
3	Construction of human activity-based phosphorylation networks. <i>Molecular Systems Biology</i> , 2013, 9, 655.	7.2	153
4	Regulation of CK2 by phosphorylation and O-GlcNAcylation revealed by semisynthesis. <i>Nature Chemical Biology</i> , 2012, 8, 262-269.	8.0	148
5	Rapid Identification of Monospecific Monoclonal Antibodies Using a Human Proteome Microarray. <i>Molecular and Cellular Proteomics</i> , 2012, 11, O111.016253.	3.8	136
6	The <i>Magnaporthe grisea</i> nod1 homolog, MSP1, is required for virulence. <i>FEMS Microbiology Letters</i> , 2007, 273, 157-165.	1.8	121
7	Identification of New Autoantigens for Primary Biliary Cirrhosis Using Human Proteome Microarrays. <i>Molecular and Cellular Proteomics</i> , 2012, 11, 669-680.	3.8	80
8	Protein Microarray Characterization of the S-Nitrosoproteome. <i>Molecular and Cellular Proteomics</i> , 2014, 13, 63-72.	3.8	56
9	Global tumor protein p53/p63 interactome. <i>Cell Cycle</i> , 2012, 11, 2367-2379.	2.6	39
10	Penetrance of Congenital Heart Disease in a Mouse Model of Down Syndrome Depends on a Trisomic Potentiator of a Disomic Modifier. <i>Genetics</i> , 2016, 203, 763-770.	2.9	31
11	Identification of SUMO E3 Ligase-Specific Substrates Using the HuProt Human Proteome Microarray. <i>Methods in Molecular Biology</i> , 2015, 1295, 455-463.	0.9	11
12	Protein Microarrays: Flexible Tools for Scientific Innovation. <i>Cold Spring Harbor Protocols</i> , 2016, 2016, pdb.top081471.	0.3	9
13	A Functional Protein Microarray Approach to Characterizing Posttranslational Modifications on Lysine Residues. <i>Methods in Molecular Biology</i> , 2011, 723, 213-223.	0.9	9
14	Posttranslational Modification Assays on Functional Protein Microarrays. <i>Cold Spring Harbor Protocols</i> , 2016, 2016, pdb.prot087999.	0.3	2