Etsuko Miyamoto-Sato

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5405802/publications.pdf

Version: 2024-02-01

20 papers 1,452 citations

687363 13 h-index 752698 20 g-index

21 all docs

21 docs citations

times ranked

21

2898 citing authors

#	Article	IF	CITATIONS
1	Six Classes of Nuclear Localization Signals Specific to Different Binding Grooves of Importin α. Journal of Biological Chemistry, 2009, 284, 478-485.	3.4	513
2	The transcriptional network that controls growth arrest and differentiation in a human myeloid leukemia cell line. Nature Genetics, 2009, 41, 553-562.	21.4	408
3	In vitro evolution of single-chain antibodies using mRNA display. Nucleic Acids Research, 2006, 34, e127-e127.	14.5	69
4	Fluorescence labeling of the C-terminus of proteins with a puromycin analogue in cell-free translation systems. FEBS Letters, 1999, 462, 43-46.	2.8	60
5	A Comprehensive Resource of Interacting Protein Regions for Refining Human Transcription Factor Networks. PLoS ONE, 2010, 5, e9289.	2.5	56
6	Highly stable and efficient mRNA templates for mRNA-protein fusions and C-terminally labeled proteins. Nucleic Acids Research, 2003, 31, 78e-78.	14.5	53
7	In vitro selection of Jun-associated proteins using mRNA display. Nucleic Acids Research, 2004, 32, e169-e169.	14.5	46
8	Cell-free cotranslation and selection using in vitro virus for high-throughput analysis of protein-protein interactions and complexes. Genome Research, 2005, 15, 710-717.	5.5	42
9	Protein–protein interaction analysis by C-terminally specific fluorescence labeling and fluorescence cross-correlation spectroscopy. Nucleic Acids Research, 2006, 34, e102-e102.	14.5	37
10	Rapid antibody selection by mRNA display on a microfluidic chip. Nucleic Acids Research, 2009, 37, e64-e64.	14.5	35
11	Next-Generation Technologies for Multiomics Approaches Including Interactome Sequencing. BioMed Research International, 2015, 2015, 1-9.	1.9	33
12	Mitochondria–Nucleus Shuttling FK506-Binding Protein 51 Interacts with TRAF Proteins and Facilitates the RIG-I-Like Receptor-Mediated Expression of Type I IFN. PLoS ONE, 2014, 9, e95992.	2.5	31
13	mRNA display selection of a highâ€affinity, Bclâ€X _L â€specific binding peptide. FASEB Journal, 2010, 24, 2201-2210.	0.5	15
14	PRD: A protein–RNA interaction database. Bioinformation, 2012, 8, 729-730.	0.5	15
15	Catalytic subunits of the phosphatase calcineurin interact with NF-κB-inducing kinase (NIK) and attenuate NIK-dependent gene expression. Scientific Reports, 2015, 5, 10758.	3.3	13
16	In vitro selection of GTP-binding proteins by block shuffling of estrogen-receptor fragments. Biochemical and Biophysical Research Communications, 2009, 390, 689-693.	2.1	10
17	Towards Personalized Medicine Mediated by in Vitro Virus-Based Interactome Approaches. International Journal of Molecular Sciences, 2014, 15, 6717-6724.	4.1	8
18	Toward functional analysis of protein interactome using "in vitrovirus―ln silicoanalyses of Fos/Jun interactors. Journal of Drug Targeting, 2006, 14, 505-511.	4.4	5

#	Article	IF	CITATIONS
19	Cell-Free Technologies for Proteomics and Protein Engineering. Protein and Peptide Letters, 2016, 23, 819-827.	0.9	2
20	Development of a novel conditional knockdown mouse based on YBâ€l protein degradation. Genes To Cells, 2018, 23, 860-867.	1.2	1