

# SAINTexpress: Improvements and additional features in INTeractome software

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Citation Report

#	ARTICLE	IF	CITATIONS
2	Protein Interaction Network of the Mammalian Hippo Pathway Reveals Mechanisms of Kinase-Phosphatase Interactions. <i>Science Signaling</i> , 2013, 6, rs15.	1.6	411
3	Spotlite: Web Application and Augmented Algorithms for Predicting Co-Complexed Proteins from Affinity Purification $\hat{=}$ Mass Spectrometry Data. <i>Journal of Proteome Research</i> , 2014, 13, 5944-5955.	1.8	18
4	CIG-P: Circular Interaction Graph for Proteomics. <i>BMC Bioinformatics</i> , 2014, 15, 344.	1.2	0
5	The interactome of the atypical phosphatase Rtr1 in <i>Saccharomyces cerevisiae</i> . <i>Molecular BioSystems</i> , 2014, 10, 1730-1741.	2.9	36
6	A Quantitative Chaperone Interaction Network Reveals the Architecture of Cellular Protein Homeostasis Pathways. <i>Cell</i> , 2014, 158, 434-448.	13.5	335
8	A Dynamic Protein Interaction Landscape of the Human Centrosome-Cilium Interface. <i>Cell</i> , 2015, 163, 1484-1499.	13.5	446
9	Rapid Analyses of Proteomes and Interactomes Using an Integrated Solid-Phase Extraction $\hat{=}$ Liquid Chromatography $\hat{=}$ MS/MS System. <i>Journal of Proteome Research</i> , 2015, 14, 977-985.	1.8	6
10	Myotubularin-related Proteins 3 and 4 Interact with Polo-like Kinase 1 and Centrosomal Protein of 55 kDa to Ensure Proper Abscission. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 946-960.	2.5	17
11	A web $\hat{=}$ tool for visualizing quantitative protein $\hat{=}$ protein interaction data. <i>Proteomics</i> , 2015, 15, 1432-1436.	1.3	30
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13	Extracting high confidence protein interactions from affinity purification data: At the crossroads. <i>Journal of Proteomics</i> , 2015, 118, 63-80.	1.2	30
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16	Identification of Regulatory and Cargo Proteins of Endosomal and Secretory Pathways in <i>Arabidopsis thaliana</i> by Proteomic Dissection*. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 1796-1813.	2.5	101
17	Applications of targeted proteomics in systems biology and translational medicine. <i>Proteomics</i> , 2015, 15, 3193-3208.	1.3	182
18	The Human Adenovirus Type 5 E4orf4 Protein Targets Two Phosphatase Regulators of the Hippo Signaling Pathway. <i>Journal of Virology</i> , 2015, 89, 8855-8870.	1.5	10
19	Biochemical Large-Scale Interaction Analysis of Murine Olfactory Receptors and Associated Signaling Proteins with Post-Synaptic Density 95, <i>Drosophila</i> Discs Large, Zona-Occludens 1 (PDZ) Domains. <i>Molecular and Cellular Proteomics</i> , 2015, 14, 2072-2084.	2.5	5
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22	From pathways to networks: Connecting dots by establishing protein-protein interaction networks in signaling pathways using affinity purification and mass spectrometry. <i>Proteomics</i> , 2015, 15, 188-202.	1.3	20
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38	Immunoprecipitation and mass spectrometry defines an extensive RBM45 protein-protein interaction network. <i>Brain Research</i> , 2016, 1647, 79-93.	1.1	43

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