

# Dã;vid Fazekas

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1247384/publications.pdf>

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13  
papers

672  
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932766

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1125271

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1310  
citing authors

#	ARTICLE	IF	CITATIONS
1	SignaLink3: a multi-layered resource to uncover tissue-specific signaling networks. <i>Nucleic Acids Research</i> , 2022, 50, D701-D709.	6.5	19
2	HSF1Base: A Comprehensive Database of HSF1 (Heat Shock Factor 1) Target Genes. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5815.	1.8	56
3	SignaLink: Multilayered Regulatory Networks. <i>Methods in Molecular Biology</i> , 2018, 1819, 53-73.	0.4	17
4	Neighbours of cancer-related proteins have key influence on pathogenesis and could increase the drug target space for anticancer therapies. <i>Npj Systems Biology and Applications</i> , 2017, 3, 2.	1.4	24
5	SalmoNet, an integrated network of ten <i>Salmonella enterica</i> strains reveals common and distinct pathways to host adaptation. <i>Npj Systems Biology and Applications</i> , 2017, 3, 31.	1.4	28
6	Identification of critical paralog groups with indispensable roles in the regulation of signaling flow. <i>Scientific Reports</i> , 2016, 6, 38588.	1.6	8
7	SignaFish: A Zebrafish-Specific Signaling Pathway Resource. <i>Zebrafish</i> , 2016, 13, 541-544.	0.5	8
8	ComPPI: a cellular compartment-specific database for protein-protein interaction network analysis. <i>Nucleic Acids Research</i> , 2015, 43, D485-D493.	6.5	116
9	SignaLink 2 – a signaling pathway resource with multi-layered regulatory networks. <i>BMC Systems Biology</i> , 2013, 7, 7.	3.0	169
10	NRF2-ome: An Integrated Web Resource to Discover Protein Interaction and Regulatory Networks of NRF2. <i>Oxidative Medicine and Cellular Longevity</i> , 2013, 2013, 1-9.	1.9	38
11	The NRF2-related interactome and regulome contain multifunctional proteins and fine-tuned autoregulatory loops. <i>FEBS Letters</i> , 2012, 586, 1795-1802.	1.3	95
12	Signalogs: Orthology-Based Identification of Novel Signaling Pathway Components in Three Metazoans. <i>PLoS ONE</i> , 2011, 6, e19240.	1.1	22
13	Uniformly curated signaling pathways reveal tissue-specific cross-talks and support drug target discovery. <i>Bioinformatics</i> , 2010, 26, 2042-2050.	1.8	72