Tim Wang

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	GCN2 adapts protein synthesis to scavenging-dependent growth. Cell Systems, 2022, 13, 158-172.e9.	2.9	12
2	Cellpose: a generalist algorithm for cellular segmentation. Nature Methods, 2021, 18, 100-106.	9.0	1,375
3	A Genome-Wide CRISPR/Cas9-Based Screen Identifies Heparan Sulfate Proteoglycans as Ligands of Killer-Cell Immunoglobulin-Like Receptors. Frontiers in Immunology, 2021, 12, 798235.	2.2	2
4	EASI-FISH for thick tissue defines lateral hypothalamus spatio-molecular organization. Cell, 2021, 184, 6361-6377.e24.	13.5	72
5	Metabolic determinants of cellular fitness dependent on mitochondrial reactive oxygen species. Science Advances, 2020, 6, .	4.7	28
6	CHP1 Regulates Compartmentalized Glycerolipid Synthesis by Activating GPAT4. Molecular Cell, 2019, 74, 45-58.e7.	4.5	83
7	Genome-Wide CRISPR/Cas9 Screening for Identification of Cancer Genes in Cell Lines. Methods in Molecular Biology, 2019, 1907, 125-136.	0.4	16
8	Paring down to the essentials. Science, 2018, 362, 904-904.	6.0	0
9	SFXN1 is a mitochondrial serine transporter required for one-carbon metabolism. Science, 2018, 362, .	6.0	154
10	Gene Essentiality Profiling Reveals Gene Networks and Synthetic Lethal Interactions with Oncogenic Ras. Cell, 2017, 168, 890-903.e15.	13.5	535
11	A genome-wide CRISPR screen identifies a restricted set of HIV host dependency factors. Nature Genetics, 2017, 49, 193-203.	9.4	290
12	A CRISPR screen identifies a pathway required for paraquat-induced cell death. Nature Chemical Biology, 2017, 13, 1274-1279.	3.9	138
13	A Genome-wide CRISPR Screen in Toxoplasma Identifies Essential Apicomplexan Genes. Cell, 2016, 166, 1423-1435.e12.	13.5	667
14	Absolute Quantification of Matrix Metabolites Reveals the Dynamics of Mitochondrial Metabolism. Cell, 2016, 166, 1324-1337.e11.	13.5	367
15	Single Guide RNA Library Design and Construction. Cold Spring Harbor Protocols, 2016, 2016, pdb.prot090803.	0.2	30
16	Viral Packaging and Cell Culture for CRISPR-Based Screens. Cold Spring Harbor Protocols, 2016, 2016, pdb.prot090811.	0.2	27
17	Large-Scale Single Guide RNA Library Construction and Use for CRISPR–Cas9-Based Genetic Screens. Cold Spring Harbor Protocols, 2016, 2016, pdb.top086892	0.2	20
18	The CASTOR Proteins Are Arginine Sensors for the mTORC1 Pathway. Cell, 2016, 165, 153-164.	13.5	598

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19	Structural basis for leucine sensing by the Sestrin2-mTORC1 pathway. Science, 2016, 351, 53-58.	6.0	340
20	Lysosomal amino acid transporter SLC38A9 signals arginine sufficiency to mTORC1. Science, 2015, 347, 188-194.	6.0	662
21	An Essential Role of the Mitochondrial Electron Transport Chain in Cell Proliferation Is to Enable Aspartate Synthesis. Cell, 2015, 162, 540-551.	13.5	1,024
22	Identification and characterization of essential genes in the human genome. Science, 2015, 350, 1096-1101.	6.0	1,461
23	Systematic identification of signaling pathways with potential to confer anticancer drug resistance. Science Signaling, 2014, 7, ra121.	1.6	163
24	Metabolic determinants of cancer cell sensitivity to glucose limitation and biguanides. Nature, 2014, 508, 108-112.	13.7	585
25	Genetic Screens in Human Cells Using the CRISPR-Cas9 System. Science, 2014, 343, 80-84.	6.0	2,414
26	Perturbation of m6A Writers Reveals Two Distinct Classes of mRNA Methylation at Internal and 5′ Sites. Cell Reports, 2014, 8, 284-296.	2.9	972
27	The Folliculin Tumor Suppressor Is a GAP for the RagC/D GTPases That Signal Amino Acid Levels to mTORC1. Molecular Cell, 2013, 52, 495-505.	4.5	436
28	MCT1-mediated transport of a toxic molecule is an effective strategy for targeting glycolytic tumors. Nature Genetics, 2013, 45, 104-108.	9.4	204