Yiwen Chen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Integrative analyses reveal a long noncoding RNA-mediated sponge regulatory network in prostate cancer. Nature Communications, 2016, 7, 10982.	12.8	267
2	Genome-wide CRISPR screen identifies HNRNPL as a prostate cancer dependency regulating RNA splicing. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E5207-E5215.	7.1	266
3	Capturing the interactome of newly transcribed RNA. Nature Methods, 2018, 15, 213-220.	19.0	170
4	The Emerging Function and Mechanism of ceRNAs in Cancer. Trends in Genetics, 2016, 32, 211-224.	6.7	164
5	Pharmacological targeting of MYC-regulated IRE1/XBP1 pathway suppresses MYC-driven breast cancer. Journal of Clinical Investigation, 2018, 128, 1283-1299.	8.2	163
6	Genome-wide identification and differential analysis of translational initiation. Nature Communications, 2017, 8, 1749.	12.8	100
7	Qki deficiency maintains stemness of glioma stem cells in suboptimal environment by downregulating endolysosomal degradation. Nature Genetics, 2017, 49, 75-86.	21.4	74
8	Mature myelin maintenance requires Qki to coactivate PPARβ-RXRα–mediated lipid metabolism. Journal of Clinical Investigation, 2020, 130, 2220-2236.	8.2	50
9	Aberrant Activation of β-Catenin Signaling Drives Glioma Tumorigenesis via USP1-Mediated Stabilization of EZH2. Cancer Research, 2019, 79, 72-85.	0.9	48
10	Exploring genetic associations with ceRNA regulation in the human genome. Nucleic Acids Research, 2017, 45, 5653-5665.	14.5	39
11	Systematic decomposition of sequence determinants governing CRISPR/Cas9 specificity. Nature Communications, 2022, 13, 474.	12.8	23
12	FGFR1-Activated Translation of WNT Pathway Components with Structured 5′ UTRs Is Vulnerable to Inhibition of EIF4A-Dependent Translation Initiation. Cancer Research, 2018, 78, 4229-4240.	0.9	22
13	Qki activates Srebp2-mediated cholesterol biosynthesis for maintenance of eye lens transparency. Nature Communications, 2021, 12, 3005.	12.8	22
14	Integrating genome-wide CRISPR immune screen with multi-omic clinical data reveals distinct classes of tumor intrinsic immune regulators. , 2021, 9, e001819.		19
15	Qki regulates myelinogenesis through Srebp2-dependent cholesterol biosynthesis. ELife, 2021, 10, .	6.0	13
16	Systematic functional interrogation of human pseudogenes using CRISPRi. Genome Biology, 2021, 22, 240.	8.8	13
17	Qki is an essential regulator of microglial phagocytosis in demyelination. Journal of Experimental Medicine, 2021, 218, .	8.5	13
18	A Proteomic Connectivity Map for Characterizing the Tumor Adaptive Response to Small Molecule Chemical Perturbagens. ACS Chemical Biology, 2020, 15, 140-150.	3.4	8

#	Article	IF	CITATIONS
19	GuidePro: a multi-source ensemble predictor for prioritizing sgRNAs in CRISPR/Cas9 protein knockouts. Bioinformatics, 2021, 37, 134-136.	4.1	7