Denis Torre

List of Publications by Year in descending order

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DENIS TOPPE

#	Article	IF	CITATIONS
1	ChEA3: transcription factor enrichment analysis by orthogonal omics integration. Nucleic Acids Research, 2019, 47, W212-W224.	14.5	512
2	Massive mining of publicly available RNA-seq data from human and mouse. Nature Communications, 2018, 9, 1366.	12.8	507
3	The Library of Integrated Network-Based Cellular Signatures NIH Program: System-Level Cataloging of Human Cells Response to Perturbations. Cell Systems, 2018, 6, 13-24.	6.2	327
4	BioJupies: Automated Generation of Interactive Notebooks for RNA-Seq Data Analysis in the Cloud. Cell Systems, 2018, 7, 556-561.e3.	6.2	217
5	Data Portal for the Library of Integrated Network-based Cellular Signatures (LINCS) program: integrated access to diverse large-scale cellular perturbation response data. Nucleic Acids Research, 2018, 46, D558-D566.	14.5	143
6	eXpression2Kinases (X2K) Web: linking expression signatures to upstream cell signaling networks. Nucleic Acids Research, 2018, 46, W171-W179.	14.5	127
7	Trained Immunity-Promoting Nanobiologic Therapy Suppresses Tumor Growth and Potentiates Checkpoint Inhibition. Cell, 2020, 183, 786-801.e19.	28.9	101
8	Toward a fine-scale population health monitoring system. Cell, 2021, 184, 2068-2083.e11.	28.9	78
9	Geneshot: search engine for ranking genes from arbitrary text queries. Nucleic Acids Research, 2019, 47, W571-W577.	14.5	57
10	KEA3: improved kinase enrichment analysis via data integration. Nucleic Acids Research, 2021, 49, W304-W316.	14.5	55
11	PPARdelta activation induces metabolic and contractile maturation of human pluripotent stem cell-derived cardiomyocytes. Cell Stem Cell, 2022, 29, 559-576.e7.	11.1	34
12	Cell-autonomous and cell non-autonomous downregulation of tumor suppressor DAB2IP by microRNA-149-3p promotes aggressiveness of cancer cells. Cell Death and Differentiation, 2018, 25, 1224-1238.	11.2	33
13	FAIRshake: Toolkit to Evaluate the FAIRness of Research Digital Resources. Cell Systems, 2019, 9, 417-421.	6.2	33
14	PRDM15 is a key regulator of metabolism critical to sustain B-cell lymphomagenesis. Nature Communications, 2020, 11, 3520.	12.8	20
15	Interoperable RNA-Seq analysis in the cloud. Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms, 2020, 1863, 194521.	1.9	17
16	Datasets2Tools, repository and search engine for bioinformatics datasets, tools and canned analyses. Scientific Data, 2018, 5, 180023.	5.3	12
17	Transcriptomic profiling of recessive dystrophic epidermolysis bullosa wounded skin highlights drug repurposing opportunities to improve wound healing. Experimental Dermatology, 2022, 31, 420-426.	2.9	9
18	Development of Potent Cellular and Humoral Immune Responses in Long-Term Hemodialysis Patients After 1273-mRNA SARS-CoV-2 Vaccination. Frontiers in Immunology, 2022, 13, 845882.	4.8	6

#	Article	IF	CITATIONS
19	Leveraging health systems data to characterize a large effect variant conferring risk for liver disease in Puerto Ricans. American Journal of Human Genetics, 2021, 108, 2099-2111.	6.2	4