## Loyal A Goff

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Differential gene and transcript expression analysis of RNA-seq experiments with TopHat and Cufflinks. Nature Protocols, 2012, 7, 562-578.	5.5	11,433
2	Differential analysis of gene regulation at transcript resolution with RNA-seq. Nature Biotechnology, 2013, 31, 46-53.	9.4	3,256
3	Integrative annotation of human large intergenic noncoding RNAs reveals global properties and specific subclasses. Genes and Development, 2011, 25, 1915-1927.	2.7	3,208
4	Multiple knockout mouse models reveal lincRNAs are required for life and brain development. ELife, 2013, 2, e01749.	2.8	609
5	Topological organization of multichromosomal regions by the long intergenic noncoding RNA Firre. Nature Structural and Molecular Biology, 2014, 21, 198-206.	3.6	565
6	DNMT1-interacting RNAs block gene-specific DNA methylation. Nature, 2013, 503, 371-376.	13.7	446
7	Long noncoding RNAs regulate adipogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 3387-3392.	3.3	371
8	Single-Cell RNA-Seq Analysis of Retinal Development Identifies NFI Factors as Regulating Mitotic Exit and Late-Born Cell Specification. Neuron, 2019, 102, 1111-1126.e5.	3.8	343
9	Linear models enable powerful differential activity analysis in massively parallel reporter assays. BMC Genomics, 2019, 20, 209.	1.2	322
10	Differential regulation of microRNA stability. Rna, 2010, 16, 1032-1039.	1.6	253
11	DeCoN: Genome-wide Analysis of InÂVivo Transcriptional Dynamics during Pyramidal Neuron Fate Selection in Neocortex. Neuron, 2015, 85, 275-288.	3.8	248
12	The long non-coding RNA Morrbid regulates Bim and short-lived myeloid cell lifespan. Nature, 2016, 537, 239-243.	13.7	234
13	The MicroRNA miR-181 Is a Critical Cellular Metabolic Rheostat Essential for NKT Cell Ontogenesis and Lymphocyte Development and Homeostasis. Immunity, 2013, 38, 984-997.	6.6	223
14	Enter the Matrix: Factorization Uncovers Knowledge from Omics. Trends in Genetics, 2018, 34, 790-805.	2.9	181
15	Transcriptional and epigenomic landscapes of CNS and non-CNS vascular endothelial cells. ELife, 2018, 7, .	2.8	180
16	MicroRNA Expression Pattern of Undifferentiated and Differentiated Human Embryonic Stem Cells. Stem Cells and Development, 2007, 16, 1003-1016.	1.1	173
17	Single-Cell Analysis of Human Retina Identifies Evolutionarily Conserved and Species-Specific Mechanisms Controlling Development. Developmental Cell, 2020, 53, 473-491.e9.	3.1	170
18	Linking RNA biology to IncRNAs. Genome Research, 2015, 25, 1456-1465.	2.4	158

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19	Spatiotemporal expression and transcriptional perturbations by long noncoding RNAs in the mouse brain. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 6855-6862.	3.3	152
20	Gene co-regulation by Fezf2 selects neurotransmitter identity and connectivity of corticospinal neurons. Nature Neuroscience, 2014, 17, 1046-1054.	7.1	121
21	Decomposing Cell Identity for Transfer Learning across Cellular Measurements, Platforms, Tissues, and Species. Cell Systems, 2019, 8, 395-411.e8.	2.9	121
22	Ago2 Immunoprecipitation Identifies Predicted MicroRNAs in Human Embryonic Stem Cells and Neural Precursors. PLoS ONE, 2009, 4, e7192.	1.1	103
23	A ketogenic diet rescues hippocampal memory defects in a mouse model of Kabuki syndrome. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 125-130.	3.3	102
24	Single-Cell RNA-Seq of Mouse Dopaminergic Neurons Informs Candidate Gene Selection for Sporadic Parkinson Disease. American Journal of Human Genetics, 2018, 102, 427-446.	2.6	102
25	Differentiating human multipotent mesenchymal stromal cells regulate microRNAs: Prediction of microRNA regulation by PDGF during osteogenesis. Experimental Hematology, 2008, 36, 1354-1369.e2.	0.2	88
26	Changes in the Excitability of Neocortical Neurons in a Mouse Model of Amyotrophic Lateral Sclerosis Are Not Specific to Corticospinal Neurons and Are Modulated by Advancing Disease. Journal of Neuroscience, 2017, 37, 9037-9053.	1.7	81
27	RNase-mediated protein footprint sequencing reveals protein-binding sites throughout the human transcriptome. Genome Biology, 2014, 15, R3.	13.9	76
28	Expression profiling of synaptic microRNAs from the adult rat brain identifies regional differences and seizure-induced dynamic modulation. Brain Research, 2012, 1436, 20-33.	1.1	68
29	Rational Probe Optimization and Enhanced Detection Strategy for MicroRNAs Using Microarrays. RNA Biology, 2005, 2, 93-100.	1.5	58
30	Group 1 Innate Lymphoid Cell Lineage Identity Is Determined by a cis-Regulatory Element Marked by a Long Non-coding RNA. Immunity, 2017, 47, 435-449.e8.	6.6	57
31	Parallel Social Information Processing Circuits Are Differentially Impacted in Autism. Neuron, 2020, 108, 659-675.e6.	3.8	52
32	Computational analysis of noncoding RNAs. Wiley Interdisciplinary Reviews RNA, 2012, 3, 759-778.	3.2	50
33	Evaluation of sense-strand mRNA amplification by comparative quantitative PCR. BMC Genomics, 2004, 5, 76.	1.2	45
34	projectR: an R/Bioconductor package for transfer learning via PCA, NMF, correlation and clustering. Bioinformatics, 2020, 36, 3592-3593.	1.8	45
35	Hypoxia tolerance in the Norrin-deficient retina and the chronically hypoxic brain studied at single-cell resolution. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 9103-9114.	3.3	44
36	Precocious neuronal differentiation and disrupted oxygen responses in Kabuki syndrome. JCI Insight, 2019, 4, .	2.3	41

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37	Long noncoding RNAs: Central to nervous system development. International Journal of Developmental Neuroscience, 2016, 55, 109-116.	0.7	34
38	Comprehensive analysis of a mouse model of spontaneous uveoretinitis using single-cell RNA sequencing. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 26734-26744.	3.3	33
39	Increased expression of anion transporter SLC26A9 delays diabetes onset in cystic fibrosis. Journal of Clinical Investigation, 2019, 130, 272-286.	3.9	33
40	A screen of 1,049 schizophrenia and 30 Alzheimer'sâ€associated variants for regulatory potential. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2020, 183, 61-73.	1.1	31
41	Mitoregulin Controls β-Oxidation in Human and Mouse Adipocytes. Stem Cell Reports, 2020, 14, 590-602.	2.3	31
42	Universal prediction of cell-cycle position using transfer learning. Genome Biology, 2022, 23, 41.	3.8	30
43	Creation and characterization of an airway epithelial cell line for stable expression of CFTR variants. Journal of Cystic Fibrosis, 2016, 15, 285-294.	0.3	28
44	A Positive Feedback Mechanism That Regulates Expression of miR-9 during Neurogenesis. PLoS ONE, 2014, 9, e94348.	1.1	28
45	Differential Variation Analysis Enables Detection of Tumor Heterogeneity Using Single-Cell RNA-Sequencing Data. Cancer Research, 2019, 79, 5102-5112.	0.4	23
46	Investigating long noncoding RNAs using animal models. Journal of Clinical Investigation, 2016, 126, 2783-2791.	3.9	23
47	Follistatin promotes LIN28B-mediated supporting cell reprogramming and hair cell regeneration in the murine cochlea. Science Advances, 2022, 8, eabj7651.	4.7	21
48	Rapid Induction of Genes Associated with Tissue Protection and Neural Development in Contused Adult Spinal Cord after Radial Glial Cell Transplantation. Journal of Neurotrauma, 2009, 26, 979-993.	1.7	15
49	Functional differentiation of a clone resembling embryonic cortical interneuron progenitors. Developmental Neurobiology, 2008, 68, 1549-1564.	1.5	13
50	Screening non-MAPT genes of the Chr17q21 H1 haplotype in Parkinson's disease. Parkinsonism and Related Disorders, 2020, 78, 138-144.	1.1	12
51	Differential Expression Levels of Sox9 in Early Neocortical Radial Glial Cells Regulate the Decision between Stem Cell Maintenance and Differentiation. Journal of Neuroscience, 2021, 41, 6969-6986.	1.7	12
52	Odorant-receptor-mediated regulation of chemosensory gene expression in the malaria mosquito Anopheles gambiae. Cell Reports, 2022, 38, 110494.	2.9	12
53	Striking heterogeneity of somatic L1 retrotransposition in single normal and cancerous gastrointestinal cells. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 32215-32222.	3.3	11
54	Postnatal Smad3 Inactivation in Murine Smooth Muscle Cells Elicits a Temporally and Regionally Distinct Transcriptional Response. Frontiers in Cardiovascular Medicine, 2022, 9, 826495.	1.1	7

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55	Poly-combing the genome for RNA. Nature Structural and Molecular Biology, 2013, 20, 1344-1346.	3.6	6
56	Bioinformatic analysis of neural stem cell differentiation. Journal of Biomolecular Techniques, 2007, 18, 205-12.	0.8	5
57	An in vivo screen of noncoding loci reveals that <i>Daedalus</i> is a gatekeeper of an Ikaros-dependent checkpoint during haematopoiesis. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	3.3	2
58	Single-Cell Analysis of Human Retina Identifies Evolutionarily Conserved and Species-Specific Mechanisms Controlling Development. SSRN Electronic Journal, 0, , .	0.4	2
59	Developmental, cellular, and behavioral phenotypes in a mouse model of congenital hypoplasia of the dentate gyrus. ELife, 2020, 9, .	2.8	2
60	The Analysis of MicroRNAs in Stem Cells. , 2008, , 141-167.		1
61	Rapid induction of genes associated with tissue protection and neural development in contused adult spinal cord after radial glial cell transplantation. Journal of Neurotrauma. 0. , 090330061141047.	1.7	1