Georg W Otto

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Oncogenic Gata1 causes stage-specific megakaryocyte differentiation delay. Haematologica, 2021, 106, 1106-1119.	3.5	8
2	Failures of nerve regeneration caused by aging or chronic denervation are rescued by restoring Schwann cell c-Jun. ELife, 2021, 10, .	6.0	63
3	Conserved properties of genetic architecture of renal and fat transcriptomes in rat models of insulin resistance. DMM Disease Models and Mechanisms, 2019, 12, .	2.4	6
4	Systems Genetics of Hepatic Metabolome Reveals Octopamine as a Target for Non-Alcoholic Fatty Liver Disease Treatment. Scientific Reports, 2019, 9, 3656.	3.3	11
5	Single-cell analysis reveals the continuum of human lympho-myeloid progenitor cells. Nature Immunology, 2018, 19, 85-97.	14.5	193
6	SCL/TAL1 cooperates with Polycomb RYBP-PRC1 to suppress alternative lineages in blood-fated cells. Nature Communications, 2018, 9, 5375.	12.8	29
7	Graded Elevation of c-Jun in Schwann Cells <i>In Vivo</i> : Gene Dosage Determines Effects on Development, Remyelination, Tumorigenesis, and Hypomyelination. Journal of Neuroscience, 2017, 37, 12297-12313.	3.6	66
8	The Calcineurin-FoxO-MuRF1 signaling pathway regulates myofibril integrity in cardiomyocytes. ELife, 2017, 6, .	6.0	33
9	Genetically distinct leukemic stem cells in human CD34â~' acute myeloid leukemia are arrested at a hemopoietic precursor-like stage. Journal of Experimental Medicine, 2016, 213, 1513-1535.	8.5	120
10	Topological analysis of metabolic networks integrating co-segregating transcriptomes and metabolomes in type 2 diabetic rat congenic series. Genome Medicine, 2016, 8, 101.	8.2	19
11	Transcriptome Profiling in Rat Inbred Strains and Experimental Cross Reveals Discrepant Genetic Architecture of Genome-Wide Gene Expression. G3: Genes, Genomes, Genetics, 2016, 6, 3671-3683.	1.8	9
12	A bi-modal function of Wnt signalling directs an FGF activity gradient to spatially regulate neuronal differentiation in the midbrain. Development (Cambridge), 2014, 141, 63-72.	2.5	30
13	Genetic Control of Differential Acetylation in Diabetic Rats. PLoS ONE, 2014, 9, e94555.	2.5	7
14	Genome Sequencing Reveals Loci under Artificial Selection that Underlie Disease Phenotypes in the Laboratory Rat. Cell, 2013, 154, 691-703.	28.9	154
15	Causes and Consequences of Chromatin Variation between Inbred Mice. PLoS Genetics, 2013, 9, e1003570.	3.5	18
16	The Light Responsive Transcriptome of the Zebrafish: Function and Regulation. PLoS ONE, 2011, 6, e17080.	2.5	90
17	Aplexone targets the HMG-CoA reductase pathway and differentially regulates arteriovenous angiogenesis. Development (Cambridge), 2011, 138, 1173-1181.	2.5	59
18	Simplet controls cell proliferation and gene transcription during zebrafish caudal fin regeneration. Developmental Biology, 2009, 325, 329-340.	2.0	45

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19	Differential gene expression as a toxicant-sensitive endpoint in zebrafish embryos and larvae. Aquatic Toxicology, 2007, 81, 355-364.	4.0	112
20	Transcriptome profiling of adult zebrafish at the late stage of chronic tuberculosis due to Mycobacterium marinum infection. Molecular Immunology, 2005, 42, 1185-1203.	2.2	129
21	Complex probes for high-throughput parallel genetic mapping of genomic mouse BAC clones. Mammalian Genome, 1998, 9, 611-616.	2.2	4