

Andreas Dahl

List of Publications by Year in descending order

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Version: 2024-02-01

131
papers

28,806
citations

61857

43
h-index

17546

121
g-index

150
all docs

150
docs citations

150
times ranked

50104
citing authors

#	ARTICLE	IF	CITATIONS
1	The variant call format and VCFtools. <i>Bioinformatics</i> , 2011, 27, 2156-2158.	1.8	11,326
2	A map of human genome variation from population-scale sequencing. <i>Nature</i> , 2010, 467, 1061-1073.	13.7	7,209
3	Mapping copy number variation by population-scale genome sequencing. <i>Nature</i> , 2011, 470, 59-65.	13.7	991
4	Modulation of Myelopoiesis Progenitors Is an Integral Component of Trained Immunity. <i>Cell</i> , 2018, 172, 147-161.e12.	13.5	702
5	Diversity of Human Copy Number Variation and Multicopy Genes. <i>Science</i> , 2010, 330, 641-646.	6.0	609
6	Demographic history and rare allele sharing among human populations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 11983-11988.	3.3	589
7	Variation in genome-wide mutation rates within and between human families. <i>Nature Genetics</i> , 2011, 43, 712-714.	9.4	525
8	Human-specific gene <i>ARHGAP11B</i> promotes basal progenitor amplification and neocortex expansion. <i>Science</i> , 2015, 347, 1465-1470.	6.0	487
9	The axolotl genome and the evolution of key tissue formation regulators. <i>Nature</i> , 2018, 554, 50-55.	13.7	463
10	The 1000 Genomes Project: data management and community access. <i>Nature Methods</i> , 2012, 9, 459-462.	9.0	308
11	Identity-by-descent filtering of exome sequence data identifies PIGV mutations in hyperphosphatasia mental retardation syndrome. <i>Nature Genetics</i> , 2010, 42, 827-829.	9.4	286
12	Innate Immune Training of Granulopoiesis Promotes Anti-tumor Activity. <i>Cell</i> , 2020, 183, 771-785.e12.	13.5	277
13	IGF2BP1 promotes SRF-dependent transcription in cancer in a m6A- and miRNA-dependent manner. <i>Nucleic Acids Research</i> , 2019, 47, 375-390.	6.5	256
14	Transcriptome sequencing during mouse brain development identifies long non-coding RNAs functionally involved in neurogenic commitment. <i>EMBO Journal</i> , 2013, 32, 3145-3160.	3.5	215
15	The genome of <i>Schmidtea mediterranea</i> and the evolution of core cellular mechanisms. <i>Nature</i> , 2018, 554, 56-61.	13.7	191
16	Defective removal of ribonucleotides from DNA promotes systemic autoimmunity. <i>Journal of Clinical Investigation</i> , 2015, 125, 413-424.	3.9	190
17	The Earliest Transcribed Zygotic Genes Are Short, Newly Evolved, and Different across Species. <i>Cell Reports</i> , 2014, 6, 285-292.	2.9	179
18	The functional spectrum of low-frequency coding variation. <i>Genome Biology</i> , 2011, 12, R84.	13.9	173

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19	Rapid Conversion of Fibroblasts into Functional Forebrain GABAergic Interneurons by Direct Genetic Reprogramming. <i>Cell Stem Cell</i> , 2015, 17, 719-734.	5.2	152
20	Reactivating head regrowth in a regeneration-deficient planarian species. <i>Nature</i> , 2013, 500, 81-84.	13.7	149
21	Mouse SAMHD1 Has Antiretroviral Activity and Suppresses a Spontaneous Cell-Intrinsic Antiviral Response. <i>Cell Reports</i> , 2013, 4, 689-696.	2.9	139
22	IL4/STAT6 Signaling Activates Neural Stem Cell Proliferation and Neurogenesis upon Amyloid- β 242 Aggregation in Adult Zebrafish Brain. <i>Cell Reports</i> , 2016, 17, 941-948.	2.9	136
23	SAMHD1 prevents autoimmunity by maintaining genome stability. <i>Annals of the Rheumatic Diseases</i> , 2015, 74, e17-e17.	0.5	133
24	Systems biology of the IMIDIA biobank from organ donors and pancreatectomised patients defines a novel transcriptomic signature of islets from individuals with type 2 diabetes. <i>Diabetologia</i> , 2018, 61, 641-657.	2.9	131
25	Genome-wide DNA Methylation Events in <i>TMPRSS2-ERG</i> Fusion-Negative Prostate Cancers Implicate an EZH2-Dependent Mechanism with <i>miR-26a</i> Hypermethylation. <i>Cancer Discovery</i> , 2012, 2, 1024-1035.	7.7	127
26	3D Culture Method for Alzheimer's Disease Modeling Reveals Interleukin-4 Rescues β 242-Induced Loss of Human Neural Stem Cell Plasticity. <i>Developmental Cell</i> , 2018, 46, 85-101.e8.	3.1	118
27	The age and genomic integrity of neurons after cortical stroke in humans. <i>Nature Neuroscience</i> , 2014, 17, 801-803.	7.1	108
28	The histone demethylase UTX regulates stem cell migration and hematopoiesis. <i>Blood</i> , 2013, 121, 2462-2473.	0.6	93
29	Single-Cell Transcriptomics Analyses of Neural Stem Cell Heterogeneity and Contextual Plasticity in a Zebrafish Brain Model of Amyloid Toxicity. <i>Cell Reports</i> , 2019, 27, 1307-1318.e3.	2.9	87
30	Multi-omics profiling of living human pancreatic islet donors reveals heterogeneous beta cell trajectories towards type 2 diabetes. <i>Nature Metabolism</i> , 2021, 3, 1017-1031.	5.1	76
31	Clonal expansion capacity defines two consecutive developmental stages of long-term hematopoietic stem cells. <i>Journal of Experimental Medicine</i> , 2014, 211, 209-215.	4.2	75
32	Insm1 Induces Neural Progenitor Delamination in Developing Neocortex via Downregulation of the Adherens Junction Belt-Specific Protein <i>Plekha7</i> . <i>Neuron</i> , 2018, 97, 1299-1314.e8.	3.8	73
33	Loss of hepatic <i>Mboat7</i> leads to liver fibrosis. <i>Gut</i> , 2021, 70, 940-950.	6.1	73
34	Effects of different management regimes on microbial biodiversity in vineyard soils. <i>Scientific Reports</i> , 2018, 8, 9393.	1.6	71
35	Dual redundant sequencing strategy: Full-length gene characterisation of 1056 novel and confirmatory <i>HLA</i> alleles. <i>Hla</i> , 2017, 90, 79-87.	0.4	70
36	Epigenomic map of human liver reveals principles of zonated morphogenic and metabolic control. <i>Nature Communications</i> , 2018, 9, 4150.	5.8	65

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37	Loss of Trex1 in Dendritic Cells Is Sufficient To Trigger Systemic Autoimmunity. <i>Journal of Immunology</i> , 2016, 197, 2157-2166.	0.4	61
38	Single cell sequencing of radial glia progeny reveals diversity of newborn neurons in the adult zebrafish brain. <i>Development (Cambridge)</i> , 2020, 147, 1855951.	1.2	60
39	A common framework of monocyte-derived macrophage activation. <i>Science Immunology</i> , 2022, 7, eabl7482.	5.6	58
40	Molecular fungal community and its decomposition activity in sapwood and heartwood of 13 temperate European tree species. <i>PLoS ONE</i> , 2019, 14, e0212120.	1.1	55
41	ROS Dynamics Delineate Functional States of Hippocampal Neural Stem Cells and Link to Their Activity-Dependent Exit from Quiescence. <i>Cell Stem Cell</i> , 2021, 28, 300-314.e6.	5.2	55
42	RNAi profiling of primary human AML cells identifies ROCK1 as a therapeutic target and nominates fasudil as an antileukemic drug. <i>Blood</i> , 2015, 125, 3760-3768.	0.6	53
43	High Diversity in the TCR Repertoire of GAD65 Autoantigen-Specific Human CD4+ T Cells. <i>Journal of Immunology</i> , 2015, 194, 2531-2538.	0.4	51
44	CCND1-CDK4-mediated cell cycle progression provides a competitive advantage for human hematopoietic stem cells in vivo. <i>Journal of Experimental Medicine</i> , 2015, 212, 1171-1183.	4.2	50
45	Multiplexing clonality: combining RGB marking and genetic barcoding. <i>Nucleic Acids Research</i> , 2014, 42, e56-e56.	6.5	49
46	An Engineered Virus Library as a Resource for the Spectrum-wide Exploration of Virus and Vector Diversity. <i>Cell Reports</i> , 2017, 19, 1698-1709.	2.9	49
47	Regulation of Liver Metabolism by the Endosomal GTPase Rab5. <i>Cell Reports</i> , 2015, 11, 884-892.	2.9	47
48	SETD1A protects HSCs from activation-induced functional decline in vivo. <i>Blood</i> , 2018, 131, 1311-1324.	0.6	47
49	Chromatoid Body Protein TDRD6 Supports Long 3' UTR Triggered Nonsense Mediated mRNA Decay. <i>PLoS Genetics</i> , 2016, 12, e1005857.	1.5	46
50	The histone 3 lysine 4 methyltransferase Setd1b is a maternal effect gene required for the oogenic gene expression program. <i>Development (Cambridge)</i> , 2017, 144, 2606-2617.	1.2	44
51	Bacteria inhabiting deadwood of 13 tree species are heterogeneously distributed between sapwood and heartwood. <i>Environmental Microbiology</i> , 2018, 20, 3744-3756.	1.8	44
52	Tox: a multifunctional transcription factor and novel regulator of mammalian corticogenesis. <i>EMBO Journal</i> , 2015, 34, 896-910.	3.5	43
53	Limitations and challenges of genetic barcode quantification. <i>Scientific Reports</i> , 2017, 7, 43249.	1.6	43
54	Odontoblast TRPC5 channels signal cold pain in teeth. <i>Science Advances</i> , 2021, 7, .	4.7	42

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55	Hsp90 inhibition differentially destabilises MAP kinase and TGF-beta signalling components in cancer cells revealed by kinase-targeted chemoproteomics. <i>BMC Cancer</i> , 2012, 12, 38.	1.1	41
56	Aldh1b1 expression defines progenitor cells in the adult pancreas and is required for Kras-induced pancreatic cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 20679-20688.	3.3	41
57	CD8+ T cells specific for the islet autoantigen IGRP are restricted in their T cell receptor chain usage. <i>Scientific Reports</i> , 2017, 7, 44661.	1.6	37
58	Quantitative PCR based expression analysis on a nanoliter scale using polymer nano-well chips. <i>Biomedical Microdevices</i> , 2007, 9, 307-314.	1.4	34
59	Donor cell leukemia: evidence for multiple preleukemic clones and parallel long term clonal evolution in donor and recipient. <i>Leukemia</i> , 2017, 31, 1637-1640.	3.3	34
60	The H3K4 methyltransferase Setd1b is essential for hematopoietic stem and progenitor cell homeostasis in mice. <i>ELife</i> , 2018, 7, .	2.8	34
61	Cre-Controlled CRISPR mutagenesis provides fast and easy conditional gene inactivation in zebrafish. <i>Nature Communications</i> , 2021, 12, 1125.	5.8	29
62	Environmental enrichment preserves a young DNA methylation landscape in the aged mouse hippocampus. <i>Nature Communications</i> , 2021, 12, 3892.	5.8	29
63	Hematopoietic Stem Cells but Not Multipotent Progenitors Drive Erythropoiesis during Chronic Erythroid Stress in EPO Transgenic Mice. <i>Stem Cell Reports</i> , 2018, 10, 1908-1919.	2.3	28
64	The complete and fully assembled genome sequence of <i>Aeromonas salmonicida</i> subsp. <i>pectinolytica</i> and its comparative analysis with other <i>Aeromonas</i> species: investigation of the mobilome in environmental and pathogenic strains. <i>BMC Genomics</i> , 2018, 19, 20.	1.2	28
65	Prospective isolation of nonhematopoietic cells of the niche and their differential molecular interactions with HSCs. <i>Blood</i> , 2019, 134, 1214-1226.	0.6	27
66	Pancreas lineage allocation and specification are regulated by sphingosine-1-phosphate signalling. <i>PLoS Biology</i> , 2017, 15, e2000949.	2.6	27
67	Abundant cytomegalovirus (CMV) reactive clonotypes in the CD8+ T cell receptor alpha repertoire following allogeneic transplantation. <i>Clinical and Experimental Immunology</i> , 2016, 184, 389-402.	1.1	26
68	Kmt2b conveys monovalent and bivalent H3K4me3 in mouse spermatogonial stem cells at germline and embryonic promoters. <i>Development (Cambridge)</i> , 2018, 145, .	1.2	26
69	Gene Expression-Based Identification of Antigen-Responsive CD8+ T Cells on a Single-Cell Level. <i>Frontiers in Immunology</i> , 2019, 10, 2568.	2.2	25
70	Continuous mitotic activity of primitive hematopoietic stem cells in adult mice. <i>Journal of Experimental Medicine</i> , 2020, 217, .	4.2	25
71	T cell receptor repertoires after adoptive transfer of expanded allogeneic regulatory T cells. <i>Clinical and Experimental Immunology</i> , 2017, 187, 316-324.	1.1	24
72	Ecophysiology of food-borne pathogens: Essential knowledge to improve food safety. <i>International Journal of Food Microbiology</i> , 2010, 139, S64-S78.	2.1	23

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73	The genome of the tegu lizard <i>Salvator merianae</i> : combining Illumina, PacBio, and optical mapping data to generate a highly contiguous assembly. <i>GigaScience</i> , 2018, 7, .	3.3	23
74	Analysis of 4 Single-Nucleotide Polymorphisms in Relation to Cervical Dysplasia and Cancer Development Using a High-Throughput Ligation-Detection Reaction Procedure. <i>International Journal of Gynecological Cancer</i> , 2011, 21, 1664-1671.	1.2	20
75	Hematopoietic stem cell response to acute thrombocytopenia requires signaling through distinct receptor tyrosine kinases. <i>Blood</i> , 2019, 134, 1046-1058.	0.6	18
76	MLL4 is required after implantation whereas MLL3 becomes essential during late gestation. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	18
77	Whole exome sequencing identifies mTOR and KEAP1 as potential targets for radiosensitization of HNSCC cells refractory to EGFR and I^21 integrin inhibition. <i>Oncotarget</i> , 2018, 9, 18099-18114.	0.8	18
78	Convergent and lineage-specific genomic differences in limb regulatory elements in limbless reptile lineages. <i>Cell Reports</i> , 2022, 38, 110280.	2.9	18
79	Aldehyde dehydrogenase activity is necessary for beta cell development and functionality in mice. <i>Diabetologia</i> , 2016, 59, 139-150.	2.9	17
80	Although Abundant in Tumor Tissue, Mast Cells Have No Effect on Immunological Micro-milieu or Growth of HPV-Induced or Transplanted Tumors. <i>Cell Reports</i> , 2018, 22, 27-35.	2.9	17
81	T-cell receptor- α repertoire of CD8+ T cells following allogeneic stem cell transplantation using next-generation sequencing. <i>Haematologica</i> , 2019, 104, 622-631.	1.7	16
82	Type 1 Interleukin-4 Signaling Obliterates Mouse Astroglia in vivo but Not in vitro. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 114.	1.8	16
83	Exosomal miRNAs from Prostate Cancer Impair Osteoblast Function in Mice. <i>International Journal of Molecular Sciences</i> , 2022, 23, 1285.	1.8	16
84	Identification and expression patterns of novel long non-coding RNAs in neural progenitors of the developing mammalian cortex. <i>Neurogenesis (Austin, Tex)</i> , 2015, 2, e995524.	1.5	15
85	TDRD6 mediates early steps of spliceosome maturation in primary spermatocytes. <i>PLoS Genetics</i> , 2017, 13, e1006660.	1.5	15
86	Neurotrophin Receptor p75NTR Regulates Immune Function of Plasmacytoid Dendritic Cells. <i>Frontiers in Immunology</i> , 2017, 8, 981.	2.2	14
87	The RNA binding protein human antigen R is a gatekeeper of liver homeostasis. <i>Hepatology</i> , 2022, 75, 881-897.	3.6	14
88	Reactive oligodendrocyte progenitor cells (re-)myelinate the regenerating zebrafish spinal cord. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	13
89	Loss of histone methyltransferase SETD1B in oogenesis results in the redistribution of genomic histone 3 lysine 4 trimethylation. <i>Nucleic Acids Research</i> , 2022, 50, 1993-2004.	6.5	13
90	A smart polymer for sequence-selective binding, pulldown, and release of DNA targets. <i>Communications Biology</i> , 2020, 3, 369.	2.0	12

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91	High-throughput Universal Probe Salmonella Serotyping (UPSS) by nanoPCR. <i>Journal of Microbiological Methods</i> , 2010, 83, 217-223.	0.7	11
92	Primary Spinal OPC Culture System from Adult Zebrafish to Study Oligodendrocyte Differentiation In Vitro. <i>Frontiers in Cellular Neuroscience</i> , 2017, 11, 284.	1.8	11
93	MicroRNA profiling of mouse cortical progenitors and neurons reveals miR-486-5p as a regulator of neurogenesis. <i>Development (Cambridge)</i> , 2020, 147, .	1.2	11
94	Zebrafish In-Vivo Screening for Compounds Amplifying Hematopoietic Stem and Progenitor Cells: - Preclinical Validation in Human CD34+ Stem and Progenitor Cells. <i>Scientific Reports</i> , 2017, 7, 12084.	1.6	10
95	Comparative RNAi Screens in Isogenic Human Stem Cells Reveal SMARCA4 as a Differential Regulator. <i>Stem Cell Reports</i> , 2019, 12, 1084-1098.	2.3	10
96	Sequence and expression levels of circular RNAs in progenitor cell types during mouse corticogenesis. <i>Life Science Alliance</i> , 2019, 2, e201900354.	1.3	10
97	Gene-selective transcription promotes the inhibition of tissue reparative macrophages by TNF. <i>Life Science Alliance</i> , 2022, 5, e202101315.	1.3	10
98	Automated solid-phase extraction for purification of single nucleotide polymorphism genotyping products prior to matrix-assisted laser desorption/ionisation time-of-flight mass spectrometric analysis. <i>Journal of Chromatography A</i> , 2004, 1049, 9-16.	1.8	9
99	Clonal Analysis Delineates Transcriptional Programs of Osteogenic and Adipogenic Lineages of Adult Mouse Skeletal Progenitors. <i>Stem Cell Reports</i> , 2018, 11, 212-227.	2.3	9
100	Exome sequencing identifies frequent genomic loss of TET1 in IDH-wild-type glioblastoma. <i>Neoplasia</i> , 2020, 22, 800-808.	2.3	9
101	Deep Learning Improves Pancreatic Cancer Diagnosis Using RNA-Based Variants. <i>Cancers</i> , 2021, 13, 2654.	1.7	9
102	Renin cells with defective Gs β /cAMP signaling contribute to renal endothelial damage. <i>Pflügers Archiv European Journal of Physiology</i> , 2019, 471, 1205-1217.	1.3	8
103	Low Threshold for Cutaneous Allergen Sensitization but No Spontaneous Dermatitis or Atopy in FLG-Deficient Mice. <i>Journal of Investigative Dermatology</i> , 2021, 141, 2611-2619.e2.	0.3	8
104	Matrix-Assisted Laser Desorption/Ionization Mass Spectrometric Analysis of DNA on Microarrays. <i>Clinical Chemistry</i> , 2006, 52, 1303-1310.	1.5	7
105	Silenced ZNF154 Is Associated with Longer Survival in Resectable Pancreatic Cancer. <i>International Journal of Molecular Sciences</i> , 2019, 20, 5437.	1.8	7
106	Controlling distinct signaling states in cultured cancer cells provides a new platform for drug discovery. <i>FASEB Journal</i> , 2019, 33, 9235-9249.	0.2	7
107	Methylation-specific ligation detection reaction (msLDR): a new approach for multiplex evaluation of methylation patterns. <i>Molecular Genetics and Genomics</i> , 2011, 286, 279-91.	1.0	6
108	Automated solid-phase extraction for purification of single nucleotide polymorphism genotyping products prior to matrix-assisted laser desorption/ionisation time-of-flight mass spectrometric analysis. <i>Journal of Chromatography A</i> , 2004, 1049, 9-16.	1.8	6

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109	Identification of Tox chromatin binding properties and downstream targets by DamID-Seq. <i>Genomics Data</i> , 2016, 7, 264-268.	1.3	5
110	The application of massively parallel sequencing technologies in diagnostics. <i>F1000 Biology Reports</i> , 2010, 2, 59.	4.0	5
111	MLL1 is required for maintenance of intestinal stem cells. <i>PLoS Genetics</i> , 2021, 17, e1009250.	1.5	5
112	Antigen-reactive regulatory T cells can be expanded in vitro with monocytes and anti-CD28 and anti-CD154 antibodies. <i>Cytotherapy</i> , 2020, 22, 629-641.	0.3	4
113	Transcriptional profile of AvrRpt2EA-mediated resistance and susceptibility response to <i>Erwinia amylovora</i> in apple. <i>Scientific Reports</i> , 2021, 11, 8685.	1.6	4
114	A Flexible Multiwell Format for Immunofluorescence Screening Microscopy of Small-Molecule Inhibitors. <i>Assay and Drug Development Technologies</i> , 2010, 8, 571-580.	0.6	3
115	Reproducibility of 10x Genomics single cell RNA sequencing method in the immune cell environment. <i>Journal of Immunological Methods</i> , 2022, 502, 113227.	0.6	3
116	Clonal competition in BcrAbl-driven leukemia: how transplantations can accelerate clonal conversion. <i>Molecular Cancer</i> , 2017, 16, 120.	7.9	2
117	Defective removal of ribonucleotides from DNA promotes systemic lupus erythematosus. <i>Pediatric Rheumatology</i> , 2015, 13, .	0.9	1
118	Assessment of the T cell receptor repertoire in long-term platelet donors by next generation sequencing. <i>British Journal of Haematology</i> , 2018, 181, 389-391.	1.2	1
119	Automated solid-phase extraction for purification of single nucleotide polymorphism genotyping products prior to matrix-assisted laser desorption/ionisation time-of-flight mass spectrometric analysis. <i>Journal of Chromatography A</i> , 2004, 1049, 9-16.	1.8	1
120	Isolation of macrophages from mouse skin wounds for single-cell RNA sequencing. <i>STAR Protocols</i> , 2022, 3, 101337.	0.5	1
121	MsLDR-creator: a web service to design msLDR assays. <i>Molecular Genetics and Genomics</i> , 2012, 287, 273-274.	1.0	0
122	PW02-026 - Low frequency variants of NLRP3 in CAPS patients. <i>Pediatric Rheumatology</i> , 2013, 11, A167.	0.9	0
123	Assessment of clonality in BcrAbl-induced leukaemia by genetic barcodes. <i>Experimental Hematology</i> , 2014, 42, S29.	0.2	0
124	Improving the design of viral barcodes for optimal bioinformatical analysis. <i>Experimental Hematology</i> , 2014, 42, S35.	0.2	0
125	Detection of low frequency variants of the NLRP3 gene in "mutation- negative" CAPS patients using massive parallel sequencing. <i>Pediatric Rheumatology</i> , 2015, 13, .	0.9	0
126	Regulatory T cell kinetics following adoptive transfer of expanded allogeneic regulatory T cells into patients with chronic graft-versus host disease. <i>Cytotherapy</i> , 2017, 19, S11.	0.3	0

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127	Understanding the role of Mboat7 in liver disease. Zeitschrift Fur Gastroenterologie, 2021, 59, .	0.2	0
128	CCND1â€“CDK4â€“mediated cell cycle progression provides a competitive advantage for human hematopoietic stem cells in vivo. Journal of Cell Biology, 2015, 210, 2102OIA144.	2.3	0
129	Analyses of clonality of BcrAbl-induced leukemia by Genetic Barcodes. , 2017, 229, .		0
130	The RNA binding protein HuR is a master regulator of hepatic-lipid homeostasis. , 2019, 57, .		0
131	Der Verlust von intestinal epitheliale SETDB1 fÃ¼hrt zu fehlender Repression endogener Retroviren, GenotoxizitÃ¤t und intestinaler EntzÃ¼ndung. Zeitschrift Fur Gastroenterologie, 2020, 58, .	0.2	0