Takaho A Endo

List of Publications by Year in descending order

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Version: 2024-02-01

59 papers

10,930 citations

35 h-index 58 g-index

60 all docs

60 docs citations

60 times ranked

17526 citing authors

#	Article	IF	Citations
1	Combined plasma metabolomic and transcriptomic analysis identify histidine as a biomarker and potential contributor in SLE pathogenesis. Rheumatology, 2023, 62, 905-913.	1.9	3
2	Establishment of mouse stem cells that can recapitulate the developmental potential of primitive endoderm. Science, 2022, 375, 574-578.	12.6	16
3	Jasmonates and Histone deacetylase 6 activate Arabidopsis genome-wide histone acetylation and methylation during the early acute stress response. BMC Biology, 2022, 20, 83.	3.8	5
4	Sox17-mediated expression of adherent molecules is required for the maintenance of undifferentiated hematopoietic cluster formation in midgestation mouse embryos. Differentiation, 2020, 115, 53-61.	1.9	7
5	Human NK cell development in hIL-7 and hIL-15 knockin NOD/SCID/IL2rgKO mice. Life Science Alliance, 2019, 2, e201800195.	2.8	41
6	Cbfl ² 2 controls differentiation of and confers homing capacity to prethymic progenitors. Journal of Experimental Medicine, 2018, 215, 595-610.	8.5	12
7	Variant PRC1 competes with retinoic acid-related signals to repress <i>Meis2</i> in distal forelimb bud. Development (Cambridge), 2018, 145, .	2.5	15
8	Alternative pathway for the development of Vα14+ NKT cells directly from CD4–CD8– thymocytes that bypasses the CD4+CD8+ stage. Nature Immunology, 2017, 18, 274-282.	14.5	55
9	Essential Roles of SATB1 in Specifying T Lymphocyte Subsets. Cell Reports, 2017, 19, 1176-1188.	6.4	82
10	Priming of lineage-specifying genes by Bcl11b is required for lineage choice in post-selection thymocytes. Nature Communications, 2017, 8, 702.	12.8	41
11	Acetate-mediated novel survival strategy against drought in plants. Nature Plants, 2017, 3, 17097.	9.3	232
12	CFC1 is a cancer stemness-regulating factor in neuroblastoma. Oncotarget, 2017, 8, 45046-45059.	1.8	7
13	PCGF6-PRC1 suppresses premature differentiation of mouse embryonic stem cells by regulating germ cell-related genes. ELife, 2017, 6, .	6.0	99
14	Mouse oocytes suppress miR-322-5p expression in ovarian granulosa cells. Journal of Reproduction and Development, 2016, 62, 393-399.	1.4	11
15	Conversion of T cells to B cells by inactivation of polycomb-mediated epigenetic suppression of the B-lineage program. Genes and Development, 2016, 30, 2475-2485.	5.9	29
16	Activation of Endogenous Retroviruses in Dnmt1 â^'/â^' ESCs Involves Disruption of SETDB1-Mediated Repression by NP95 Binding to Hemimethylated DNA. Cell Stem Cell, 2016, 19, 81-94.	11.1	77
17	A Histone Methyltransferase ESET Is Critical for T Cell Development. Journal of Immunology, 2016, 197, 2269-2279.	0.8	33
18	Efficient Regeneration of Human Vα24+ Invariant Natural Killer T Cells and Their Anti-Tumor Activity In Vivo. Stem Cells, 2016, 34, 2852-2860.	3.2	65

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19	A Stress-Activated Transposon in Arabidopsis Induces Transgenerational Abscisic Acid Insensitivity. Scientific Reports, 2016, 6, 23181.	3.3	106
20	RING1 contributes to early proximal-distal specification of the forelimb bud by restricting Meis2 expression. Development (Cambridge), 2015, 143, 276-85.	2.5	15
21	Quality control method for RNA â€seq using single nucleotide polymorphism allele frequency. Genes To Cells, 2014, 19, 821-829.	1.2	5
22	The epigenetic regulator Uhrf1 facilitates the proliferation and maturation of colonic regulatory T cells. Nature Immunology, 2014, 15, 571-579.	14.5	147
23	Polycomb Potentiates Meis2 Activation in Midbrain by Mediating Interaction of the Promoter with a Tissue-Specific Enhancer. Developmental Cell, 2014, 28, 94-101.	7.0	96
24	KLRG ⁺ invariant natural killer T cells are long-lived effectors. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 12474-12479.	7.1	46
25	Pitfalls in global normalization of ChIP-seq data in CD4+ T cells treated with butyrate: A possible solution strategy. Genomics Data, 2014, 2, 176-180.	1.3	3
26	Basophil-Derived Interleukin-4 Controls the Function of Natural Helper Cells, a Member of ILC2s, in Lung Inflammation. Immunity, 2014, 40, 758-771.	14.3	263
27	Highly Reproducible ChIP-on-Chip Analysis to Identify Genome-Wide Protein Binding and Chromatin Status in Arabidopsis thaliana. Methods in Molecular Biology, 2014, 1062, 405-426.	0.9	6
28	SAM Domain Polymerization Links Subnuclear Clustering of PRC1 to Gene Silencing. Developmental Cell, 2013, 26, 565-577.	7.0	271
29	Commensal microbe-derived butyrate induces the differentiation of colonic regulatory T cells. Nature, 2013, 504, 446-450.	27.8	3,901
30	Embracing change to remain the same: conservation of polycomb functions despite divergence of binding motifs among species. Current Opinion in Cell Biology, 2013, 25, 305-313.	5.4	8
31	Scratch regulates neuronal migration onset via an epithelial-mesenchymal transition–like mechanism. Nature Neuroscience, 2013, 16, 416-425.	14.8	116
32	Ash1l Methylates Lys36 of Histone H3 Independently of Transcriptional Elongation to Counteract Polycomb Silencing. PLoS Genetics, 2013, 9, e1003897.	3.5	69
33	Role of SOX17 in hematopoietic development from human embryonic stem cells. Blood, 2013, 121, 447-458.	1.4	87
34	Cell Cycle-Dependent Turnover of 5-Hydroxymethyl Cytosine in Mouse Embryonic Stem Cells. PLoS ONE, 2013, 8, e82961.	2.5	73
35	Histone H2A Mono-Ubiquitination Is a Crucial Step to Mediate PRC1-Dependent Repression of Developmental Genes to Maintain ES Cell Identity. PLoS Genetics, 2012, 8, e1002774.	3.5	233
36	RYBP Represses Endogenous Retroviruses and Preimplantation- and Germ Line-Specific Genes in Mouse Embryonic Stem Cells. Molecular and Cellular Biology, 2012, 32, 1139-1149.	2.3	84

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37	Tu1922 Genome-Wide Epigenomic Analysis of Lamina Propria Mononuclear Cells of Inflammatory Bowel Disease. Gastroenterology, 2012, 142, S-878-S-879.	1.3	0
38	Genome-wide analysis of target genes regulated by HoxB4 in hematopoietic stem and progenitor cells developing from embryonic stem cells. Blood, 2011, 117, e142-e150.	1.4	42
39	The Hbo1-Brd1/Brpf2 complex is responsible for global acetylation of H3K14 and required for fetal liver erythropoiesis. Blood, 2011, 118, 2443-2453.	1.4	168
40	Mammalian Polycomb-Like Pcl2/Mtf2 Is a Novel Regulatory Component of PRC2 That Can Differentially Modulate Polycomb Activity both at the <i>Hox</i> Gene Cluster and at <i>Cdkn2a</i> Genes. Molecular and Cellular Biology, 2011, 31, 351-364.	2.3	68
41	Arabidopsis HDA6 Regulates Locus-Directed Heterochromatin Silencing in Cooperation with MET1. PLoS Genetics, 2011, 7, e1002055.	3.5	148
42	Arabidopsis Tiling Array Analysis to Identify the Stress-Responsive Genes. Methods in Molecular Biology, 2010, 639, 141-155.	0.9	27
43	Genome-wide analysis of endogenous abscisic acid-mediated transcription in dry and imbibed seeds of Arabidopsis using tiling arrays. Plant Journal, 2010, 62, 39-51.	5.7	109
44	Divergence of CpG island promoters: A consequence or cause of evolution?. Development Growth and Differentiation, 2010, 52, 545-554.	1.5	41
45	O12. Polycomb-dependent regulation for differentiation programs of stem cells and progenitors. Differentiation, 2010, 80, S9-S10.	1.9	1
46	Murine induced pluripotent stem cells can be derived from and differentiate into natural killer T cells. Journal of Clinical Investigation, 2010, 120, 2610-2618.	8.2	55
47	OmicBrowse: a Flash-based high-performance graphics interface for genomic resources. Nucleic Acids Research, 2009, 37, W57-W62.	14.5	4
48	Maintenance of Undifferentiated State and Self-Renewal of Embryonic Neural Stem Cells by Polycomb Protein Ring1B. Stem Cells, 2009, 27, 1559-1570.	3.2	57
49	Polycomb Limits the Neurogenic Competence of Neural Precursor Cells to Promote Astrogenic Fate Transition. Neuron, 2009, 63, 600-613.	8.1	420
50	Detecting Linkage between a Trait and a Marker in a Random Mating Population without Pedigree Record. PLoS ONE, 2009, 4, e4956.	2.5	0
51	Arabidopsis Transcriptome Analysis under Drought, Cold, High-Salinity and ABA Treatment Conditions using a Tiling Array. Plant and Cell Physiology, 2008, 49, 1135-1149.	3.1	475
52	Polycomb group proteins Ring1A/B are functionally linked to the core transcriptional regulatory circuitry to maintain ES cell identity. Development (Cambridge), 2008, 135, 1513-1524.	2.5	265
53	The SRA protein Np95 mediates epigenetic inheritance by recruiting Dnmt1 to methylated DNA. Nature, 2007, 450, 908-912.	27.8	1,096
54	Whole genome association study of rheumatoid arthritis using 27â€039 microsatellites. Human Molecular Genetics, 2005, 14, 2305-2321.	2.9	122

Takaho A Endo

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55	Probabilistic nucleotide assembling method for sequencing by hybridization. Bioinformatics, 2004, 20, 2181-2188.	4.1	20
56	CIS1 Interacts with the Y532 of the Prolactin Receptor and Suppresses Prolactin-Dependent STAT5 Activation. Journal of Biochemistry, 2003, 133, 109-113.	1.7	28
57	STAT5 Activation Correlates with Erythropoietin Receptor-mediated Erythroid Differentiation of an Erythroleukemia Cell Line. Journal of Biological Chemistry, 1997, 272, 8149-8152.	3.4	84
58	A new protein containing an SH2 domain that inhibits JAK kinases. Nature, 1997, 387, 921-924.	27.8	1,319
59	A Chinese Hamster Ovary Cell Mutant Resistant to Phosphatidylserine Is Defective in Transbilayer Movement of Cell Surface Phosphatidylserine. Experimental Cell Research, 1996, 228, 341-346.	2.6	12