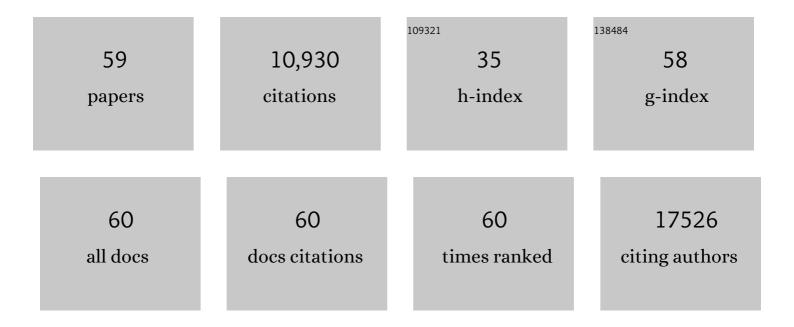
Takaho A Endo

List of Publications by Year in descending order

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Τλκλήο Δ Ενίοο

#	Article	IF	CITATIONS
1	Commensal microbe-derived butyrate induces the differentiation of colonic regulatory T cells. Nature, 2013, 504, 446-450.	27.8	3,901
2	A new protein containing an SH2 domain that inhibits JAK kinases. Nature, 1997, 387, 921-924.	27.8	1,319
3	The SRA protein Np95 mediates epigenetic inheritance by recruiting Dnmt1 to methylated DNA. Nature, 2007, 450, 908-912.	27.8	1,096
4	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
5	Polycomb Limits the Neurogenic Competence of Neural Precursor Cells to Promote Astrogenic Fate Transition. Neuron, 2009, 63, 600-613.	8.1	420
6	SAM Domain Polymerization Links Subnuclear Clustering of PRC1 to Gene Silencing. Developmental Cell, 2013, 26, 565-577.	7.0	271
7	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
8	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
9	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
10	Acetate-mediated novel survival strategy against drought in plants. Nature Plants, 2017, 3, 17097.	9.3	232
11	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
12	Arabidopsis HDA6 Regulates Locus-Directed Heterochromatin Silencing in Cooperation with MET1. PLoS Genetics, 2011, 7, e1002055.	3.5	148
13	The epigenetic regulator Uhrf1 facilitates the proliferation and maturation of colonic regulatory T cells. Nature Immunology, 2014, 15, 571-579.	14.5	147
14	Whole genome association study of rheumatoid arthritis using 27â€039 microsatellites. Human Molecular Genetics, 2005, 14, 2305-2321.	2.9	122
15	Scratch regulates neuronal migration onset via an epithelial-mesenchymal transition–like mechanism. Nature Neuroscience, 2013, 16, 416-425.	14.8	116
16	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
17	A Stress-Activated Transposon in Arabidopsis Induces Transgenerational Abscisic Acid Insensitivity. Scientific Reports, 2016, 6, 23181.	3.3	106
18	PCGF6-PRC1 suppresses premature differentiation of mouse embryonic stem cells by regulating germ cell-related genes. ELife, 2017, 6, .	6.0	99

Τακαμό Α Ενδό

#	Article	IF	CITATIONS
19	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
20	Role of SOX17 in hematopoietic development from human embryonic stem cells. Blood, 2013, 121, 447-458.	1.4	87
21	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
22	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
23	Essential Roles of SATB1 in Specifying T Lymphocyte Subsets. Cell Reports, 2017, 19, 1176-1188.	6.4	82
24	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
25	Cell Cycle-Dependent Turnover of 5-Hydroxymethyl Cytosine in Mouse Embryonic Stem Cells. PLoS ONE, 2013, 8, e82961.	2.5	73
26	Ash1l Methylates Lys36 of Histone H3 Independently of Transcriptional Elongation to Counteract Polycomb Silencing. PLoS Genetics, 2013, 9, e1003897.	3.5	69
27	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
28	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
29	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
30	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
31	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
32	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
33	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
34	Divergence of CpG island promoters: A consequence or cause of evolution?. Development Growth and Differentiation, 2010, 52, 545-554.	1.5	41
35	Priming of lineage-specifying genes by Bcl11b is required for lineage choice in post-selection thymocytes. Nature Communications, 2017, 8, 702.	12.8	41
36	Human NK cell development in hIL-7 and hIL-15 knockin NOD/SCID/IL2rgKO mice. Life Science Alliance, 2019, 2, e201800195.	2.8	41

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#	Article	IF	CITATIONS
37	A Histone Methyltransferase ESET Is Critical for T Cell Development. Journal of Immunology, 2016, 197, 2269-2279.	0.8	33
38	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
39	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
40	Arabidopsis Tiling Array Analysis to Identify the Stress-Responsive Genes. Methods in Molecular Biology, 2010, 639, 141-155.	0.9	27
41	Probabilistic nucleotide assembling method for sequencing by hybridization. Bioinformatics, 2004, 20, 2181-2188.	4.1	20
42	Establishment of mouse stem cells that can recapitulate the developmental potential of primitive endoderm. Science, 2022, 375, 574-578.	12.6	16
43	RING1 contributes to early proximal-distal specification of the forelimb bud by restricting Meis2 expression. Development (Cambridge), 2015, 143, 276-85.	2.5	15
44	Variant PRC1 competes with retinoic acid-related signals to repress <i>Meis2</i> in distal forelimb bud. Development (Cambridge), 2018, 145, .	2.5	15
45	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
46	Cbfβ2 controls differentiation of and confers homing capacity to prethymic progenitors. Journal of Experimental Medicine, 2018, 215, 595-610.	8.5	12
47	Mouse oocytes suppress miR-322-5p expression in ovarian granulosa cells. Journal of Reproduction and Development, 2016, 62, 393-399.	1.4	11
48	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
49	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
50	CFC1 is a cancer stemness-regulating factor in neuroblastoma. Oncotarget, 2017, 8, 45046-45059.	1.8	7
51	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
52	Quality control method for RNA â€seq using single nucleotide polymorphism allele frequency. Genes To Cells, 2014, 19, 821-829.	1.2	5
53	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
54	OmicBrowse: a Flash-based high-performance graphics interface for genomic resources. Nucleic Acids Research, 2009, 37, W57-W62.	14.5	4

Τακαμό Α Ενδό

#	Article	IF	CITATIONS
55	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
56	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
57	O12. Polycomb-dependent regulation for differentiation programs of stem cells and progenitors. Differentiation, 2010, 80, S9-S10.	1.9	1
58	Tu1922 Genome-Wide Epigenomic Analysis of Lamina Propria Mononuclear Cells of Inflammatory Bowel Disease. Gastroenterology, 2012, 142, S-878-S-879.	1.3	0
59	Detecting Linkage between a Trait and a Marker in a Random Mating Population without Pedigree Record. PLoS ONE, 2009, 4, e4956.	2.5	0