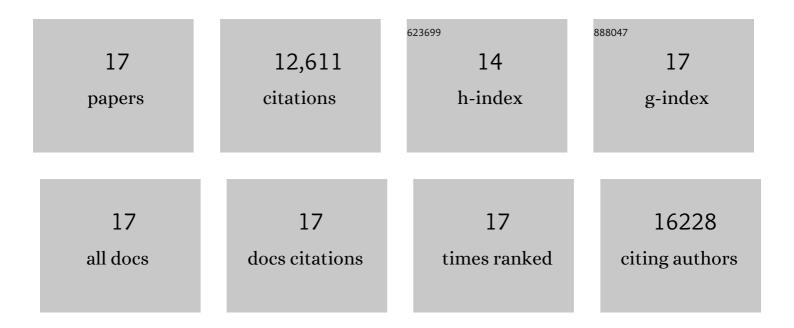
Kazunori Waki

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	The map-based sequence of the rice genome. Nature, 2005, 436, 793-800.	27.8	3,365
2	The Transcriptional Landscape of the Mammalian Genome. Science, 2005, 309, 1559-1563.	12.6	3,227
3	Antisense Transcription in the Mammalian Transcriptome. Science, 2005, 309, 1564-1566.	12.6	1,553
4	Analysis of the mouse transcriptome based on functional annotation of 60,770 full-length cDNAs. Nature, 2002, 420, 563-573.	27.8	1,548
5	The regulated retrotransposon transcriptome of mammalian cells. Nature Genetics, 2009, 41, 563-571.	21.4	731
6	Cap analysis gene expression for high-throughput analysis of transcriptional starting point and identification of promoter usage. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 15776-15781.	7.1	673
7	The genome sequence and structure of rice chromosome 1. Nature, 2002, 420, 312-316.	27.8	519
8	The transcriptional network that controls growth arrest and differentiation in a human myeloid leukemia cell line. Nature Genetics, 2009, 41, 553-562.	21.4	408
9	A Comprehensive Rice Transcript Map Containing 6591 Expressed Sequence Tag Sites. Plant Cell, 2002, 14, 525-535.	6.6	260
10	Targeting a Complex Transcriptome: The Construction of the Mouse Full-Length cDNA Encyclopedia. Genome Research, 2003, 13, 1273-1289.	5.5	154
11	Cytoskeletal Rearrangements in Synovial Fibroblasts as a Novel Pathophysiological Determinant of Modeled Rheumatoid Arthritis. PLoS Genetics, 2005, 1, e48.	3.5	49
12	Libraries enriched for alternatively spliced exons reveal splicing patterns in melanocytes and melanomas. Nature Methods, 2004, 1, 233-239.	19.0	45
13	Introduction to the Japan Mouse Clinic at the RIKEN BioResource Center. Experimental Animals, 2009, 58, 443-450.	1.1	30
14	The RIKEN integrated database of mammals. Nucleic Acids Research, 2011, 39, D861-D870.	14.5	23
15	Subtraction of cap-trapped full-length cDNA libraries to select rare transcripts. BioTechniques, 2003, 35, 510-518.	1.8	12
16	A Resource for Transcriptomic Analysis in the Mouse Brain. PLoS ONE, 2008, 3, e3012.	2.5	11
17	SDOP-DB: a comparative standardized-protocol database for mouse phenotypic analyses. Bioinformatics, 2010, 26, 1133-1134.	4.1	3