

Miyako Yamamoto

List of Publications by Year in descending order

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33
papers

1,355
citations

361413

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h-index

395702

33
g-index

33
all docs

33
docs citations

33
times ranked

968
citing authors

#	ARTICLE	IF	CITATIONS
1	ABO Research in the Modern Era of Genomics. <i>Transfusion Medicine Reviews</i> , 2012, 26, 103-118.	2.0	129
2	Molecular Genetic Analysis of the ABO Blood Group System: 2. <i>cis</i> -AB Alleles. <i>Vox Sanguinis</i> , 1993, 64, 120-123.	1.5	101
3	Molecular Genetic Analysis of the ABO Blood Group System: 4. Another Type of O Allele. <i>Vox Sanguinis</i> , 1993, 64, 175-178.	1.5	95
4	Molecular Genetic Analysis of the ABO Blood Group System: 2. <i>cis</i> -AB Alleles. <i>Vox Sanguinis</i> , 1993, 64, 120-123.	1.5	93
5	Molecular Genetic Analysis of the ABO Blood Group System: 1. Weak Subgroups: A ³ and B ³ Alleles. <i>Vox Sanguinis</i> , 1993, 64, 116-119.	1.5	92
6	Molecular Genetic Analysis of the ABO Blood Group System: 4. Another Type of O Allele. <i>Vox Sanguinis</i> , 1993, 64, 175-178.	1.5	85
7	Molecular Genetic Analysis of the ABO Blood Group System: 3. A ^x and B ^(A) Alleles. <i>Vox Sanguinis</i> , 1993, 64, 171-174.	1.5	80
8	Molecular Genetic Analysis of the ABO Blood Group System: 1. Weak Subgroups: A ³ and B ³ Alleles. <i>Vox Sanguinis</i> , 1993, 64, 116-119.	1.5	78
9	Murine Equivalent of the Human Histo-blood Group ABO Gene Is <i>cis</i> -AB Gene and Encodes a Glycosyltransferase with Both A and B Transferase Activity. <i>Journal of Biological Chemistry</i> , 2001, 276, 13701-13708.	3.4	71
10	Molecular Genetic Analysis of the ABO Blood Group System: 3. A ^x and B ^(A) Alleles. <i>Vox Sanguinis</i> , 1993, 64, 171-174.	1.5	70
11	Animal Histo-blood group ABO genes. <i>Biochemical and Biophysical Research Communications</i> , 1992, 189, 154-164.	2.1	60
12	An integrative evolution theory of histo-blood group ABO and related genes. <i>Scientific Reports</i> , 2014, 4, 6601.	3.3	48
13	Molecular genetic basis of porcine histo-blood group AO system. <i>Blood</i> , 2001, 97, 3308-3310.	1.4	42
14	NotI-MseI methylation-sensitive amplified fragment length polymorphism for DNA methylation analysis of human cancers. <i>Electrophoresis</i> , 2001, 22, 1946-1956.	2.4	40
15	Rare and Frequent Promoter Methylation, Respectively, of TSHZ2 and 3 Genes That Are Both Downregulated in Expression in Breast and Prostate Cancers. <i>PLoS ONE</i> , 2011, 6, e17149.	2.5	38
16	Molecular genetic basis of the human Forssman glycolipid antigen negativity. <i>Scientific Reports</i> , 2012, 2, 975.	3.3	33
17	Comprehensive Expression Profiling of Highly Homologous 39 Hox Genes in 26 Different Human Adult Tissues by the Modified Systematic Multiplex RT-PCR Method Reveals Tissue-Specific Expression Pattern That Suggests an Important Role of Chromosomal Structure in the Regulation of Hox Gene Expression in Adult Tissues. <i>Gene Expression</i> . 2003, 11, 199-210.	1.2	30
18	Phage Display cDNA Cloning of Protein with Carbohydrate Affinity. <i>Biochemical and Biophysical Research Communications</i> , 1999, 255, 194-199.	2.1	28

#	ARTICLE	IF	CITATIONS
19	Scanning copy number and gene expression on the 18q21-qter chromosomal region by the systematic multiplex PCR and reverse transcription-PCR methods. <i>Electrophoresis</i> , 2007, 28, 1882-1895.	2.4	23
20	Expression profiling of 68 glycosyltransferase genes in 27 different human tissues by the systematic multiplex reverse transcription-polymerase chain reaction method revealed clustering of sexually related tissues in hierarchical clustering algorithm analysis. <i>Electrophoresis</i> , 2003, 24, 2295-2307.	2.4	20
21	Crosstalk between ABO and Forssman (FORS) blood group systems: FORS1 antigen synthesis by ABO gene-encoded glycosyltransferases. <i>Scientific Reports</i> , 2017, 7, 41632.	3.3	17
22	Identification of Genes That Exhibit Changes in Expression on the 8p Chromosomal Arm by the Systematic Multiplex RT-PCR (SM RT-PCR) and DNA Microarray Hybridization Methods. <i>Gene Expression</i> , 2008, 14, 217-227.	1.2	13
23	Non-AUG start codons responsible for ABO weak blood group alleles on initiation mutant backgrounds. <i>Scientific Reports</i> , 2017, 7, 41720.	3.3	11
24	ABO blood group A transferases catalyze the biosynthesis of FORS blood group FORS1 antigen upon deletion of exon 3 or 4. <i>Blood Advances</i> , 2017, 1, 2756-2766.	5.2	9
25	Blood group ABO gene-encoded A transferase catalyzes the biosynthesis of FORS1 antigen of FORS system upon Met69Thr/Ser substitution. <i>Blood Advances</i> , 2018, 2, 1371-1381.	5.2	8
26	Systematic multiplex polymerase chain reaction and reverse transcription-polymerase chain reaction analyses of changes in copy number and expression of proto-oncogenes and tumor suppressor genes in cancer tissues and cell lines. <i>Electrophoresis</i> , 2004, 25, 3349-3356.	2.4	7
27	ABO blood group A transferase and its codon 69 substitution enzymes synthesize FORS1 antigen of FORS blood group system. <i>Scientific Reports</i> , 2019, 9, 9717.	3.3	7
28	Gene expression analysis of an integrin family of genes by systematic multiplex reverse transcription-polymerase chain reaction. <i>Electrophoresis</i> , 2004, 25, 2201-2211.	2.4	6
29	Murine Cell Glycolipids Customization by Modular Expression of Glycosyltransferases. <i>PLoS ONE</i> , 2013, 8, e64728.	2.5	6
30	IMMUNOHEMATOLOGY: Generation of histo-blood group B transferase by replacing the acetylgalactosamine recognition domain of human A transferase with the galactose recognition domain of evolutionarily related murine β 1,3-galactosyltransferase. <i>Transfusion</i> , 2010, 50, 622-630.	1.6	5
31	Amino acid substitutions at sugar-recognizing codons confer ABO blood group system-related β 1,3-Gal(NAc) transferases with differential enzymatic activity. <i>Scientific Reports</i> , 2019, 9, 846.	3.3	5
32	Scanning copy number and gene expression on the 16p13.3-13.2 chromosomal region by the systematic multiplex polymerase chain reaction and reverse transcription-polymerase chain reaction methods. <i>Electrophoresis</i> , 2006, 27, 2529-2540.	2.4	3
33	Mixed-Up Sugars: Glycosyltransferase Cross-Reactivity in Cancerous Tissues and Their Therapeutic Targeting. <i>ChemBioChem</i> , 2022, 23, .	2.6	2