

Martin J Maiers

List of Publications by Year in descending order

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Version: 2024-02-01

125
papers

10,416
citations

81889

39
h-index

34984

98
g-index

143
all docs

143
docs citations

143
times ranked

8231
citing authors

#	ARTICLE	IF	CITATIONS
1	Nomenclature for factors of the HLA system, 2010. <i>Tissue Antigens</i> , 2010, 75, 291-455.	1.0	3,121
2	HLA Match Likelihoods for Hematopoietic Stem-Cell Grafts in the U.S. Registry. <i>New England Journal of Medicine</i> , 2014, 371, 339-348.	27.0	861
3	The Shaping of Modern Human Immune Systems by Multiregional Admixture with Archaic Humans. <i>Science</i> , 2011, 334, 89-94.	12.6	441
4	High-resolution HLA alleles and haplotypes in the United States population. <i>Human Immunology</i> , 2007, 68, 779-788.	2.4	417
5	Six-locus high resolution HLA haplotype frequencies derived from mixed-resolution DNA typing for the entire US donor registry. <i>Human Immunology</i> , 2013, 74, 1313-1320.	2.4	349
6	The effect of donor characteristics on survival after unrelated donor transplantation for hematologic malignancy. <i>Blood</i> , 2016, 127, 260-267.	1.4	245
7	The Effect of KIR Ligand Incompatibility on the Outcome of Unrelated Donor Transplantation: A Report from the Center for International Blood and Marrow Transplant Research, the European Blood and Marrow Transplant Registry, and the Dutch Registry. <i>Biology of Blood and Marrow Transplantation</i> , 2006, 12, 876-884.	2.0	241
8	Impact of allele-level HLA matching on outcomes after myeloablative single unit umbilical cord blood transplantation for hematologic malignancy. <i>Blood</i> , 2014, 123, 133-140.	1.4	239
9	Progress toward an efficient panel of SNPs for ancestry inference. <i>Forensic Science International: Genetics</i> , 2014, 10, 23-32.	3.1	211
10	Selection of unrelated donors and cord blood units for hematopoietic cell transplantation: guidelines from the NMDP/CIBMTR. <i>Blood</i> , 2019, 134, 924-934.	1.4	199
11	Common and well-documented <sc>HLA</sc> alleles: 2012 update to the <sc>CWD</sc> catalogue. <i>Tissue Antigens</i> , 2013, 81, 194-203.	1.0	198
12	Common and Well-Documented HLA Alleles. <i>Human Immunology</i> , 2007, 68, 392-417.	2.4	194
13	Classification of HLA-Matching for Retrospective Analysis of Unrelated Donor Transplantation: Revised Definitions to Predict Survival. <i>Biology of Blood and Marrow Transplantation</i> , 2008, 14, 748-758.	2.0	186
14	The HLA dictionary 2008: a summary of HLAâ€A, â€B, â€C, â€DRB1/3/4/5, and â€DQB1 alleles and their association with serologically defined HLAâ€A, â€B, â€C, â€DR, and â€DQ antigens. <i>Tissue Antigens</i> , 2009, 73, 95-170.	1.0	184
15	HLA Diversity in the 1000 Genomes Dataset. <i>PLoS ONE</i> , 2014, 9, e97282.	2.5	179
16	New HLA haplotype frequency reference standards: High-resolution and large sample typing of HLA DRâ€DQ haplotypes in a sample of European Americans. <i>Tissue Antigens</i> , 2003, 62, 296-307.	1.0	157
17	Development of an Unrelated Donor Selection Score Predictive of Survival after HCT: Donor Age Matters Most. <i>Biology of Blood and Marrow Transplantation</i> , 2018, 24, 1049-1056.	2.0	98
18	The Impact of Amino Acid Variability on Alloreactivity Defines a Functional Distance Predictive of Permissive HLA-DPB1 Mismatches in Hematopoietic Stem Cell Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2015, 21, 233-241.	2.0	95

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19	Advances in the Selection of HLA-Compatible Donors: Refinements in HLA Typing and Matching over the First 20 Years of the National Marrow Donor Program Registry. <i>Biology of Blood and Marrow Transplantation</i> , 2008, 14, 37-44.	2.0	91
20	Revealing complete complex KIR haplotypes phased by long-read sequencing technology. <i>Genes and Immunity</i> , 2017, 18, 127-134.	4.1	89
21	Tracking human migrations by the analysis of the distribution of HLA alleles, lineages and haplotypes in closed and open populations. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2012, 367, 820-829.	4.0	86
22	Allele-Level Haplotype Frequencies and Pairwise Linkage Disequilibrium for 14 KIR Loci in 506 European-American Individuals. <i>PLoS ONE</i> , 2012, 7, e47491.	2.5	85
23	Availability of unrelated donors for hematopoietic stem cell transplantation for hemoglobinopathies. <i>Bone Marrow Transplantation</i> , 2003, 31, 547-550.	2.4	82
24	Amino acid substitution at peptide-binding pockets of HLA class I molecules increases risk of severe acute GVHD and mortality. <i>Blood</i> , 2013, 122, 3651-3658.	1.4	77
25	Estimation of HLA-A, -B, -DRB1 Haplotype Frequencies Using Mixed Resolution Data from a National Registry with Selective Retyping of Volunteers. <i>Human Immunology</i> , 2007, 68, 950-958.	2.4	74
26	On Modeling Human Leukocyte Antigenâ€“Identical Sibling Match Probability for Allogeneic Hematopoietic Cell Transplantation: Estimating the Need for an Unrelated Donor Source. <i>Biology of Blood and Marrow Transplantation</i> , 2016, 22, 410-417.	2.0	65
27	The HLA dictionary 1999: a summary of HLA-A, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens. <i>Tissue Antigens</i> , 1999, 54, 409-437.	1.0	63
28	HapLogic: A Predictive Human Leukocyte Antigenâ€“Matching Algorithm to Enhance Rapid Identification of the Optimal Unrelated Hematopoietic Stem Cell Sources for Transplantation. <i>Biology of Blood and Marrow Transplantation</i> , 2016, 22, 2038-2046.	2.0	63
29	Genotype List String: a grammar for describing <scp>HLA</scp> and <scp>KIR</scp> genotyping results in a text string. <i>Tissue Antigens</i> , 2013, 82, 106-112.	1.0	56
30	Genetic risk variants in African Americans with multiple sclerosis. <i>Neurology</i> , 2013, 81, 219-227.	1.1	54
31	Role of HLA-B exon 1 in graft-versus-host disease after unrelated haemopoietic cell transplantation: a retrospective cohort study. <i>Lancet Haematology</i> , 2020, 7, e50-e60.	4.6	53
32	The distribution of HLA haplotypes in the ethnic groups that make up the Brazilian Bone Marrow Volunteer Donor Registry (REDOME). <i>Immunogenetics</i> , 2018, 70, 511-522.	2.4	51
33	A community standard for immunogenomic data reporting and analysis: proposal for a STrengthening the REporting of Immunogenomic Studies statement. <i>Tissue Antigens</i> , 2011, 78, 333-344.	1.0	50
34	An update to HLA Nomenclature, 2010. <i>Bone Marrow Transplantation</i> , 2010, 45, 846-848.	2.4	48
35	Validation of statistical imputation of alleleâ€“level multilocus phased genotypes from ambiguous <scp>HLA</scp> assignments. <i>Tissue Antigens</i> , 2014, 84, 285-292.	1.0	48
36	A combined DPA1âˆ¼DPB1 amino acid epitope is the primary unit of selection on the HLA-DP heterodimer. <i>Immunogenetics</i> , 2012, 64, 559-569.	2.4	47

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37	8/8 and 10/10 High-Resolution Match Rate for the Be The Match Unrelated Donor Registry. <i>Biology of Blood and Marrow Transplantation</i> , 2015, 21, 137-141.	2.0	47
38	The HLA Dictionary 2004: a summary of HLA-A, -B, -C, -DRB1/3/4/5 and -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens. <i>Tissue Antigens</i> , 2005, 65, 1-55.	1.0	43
39	HLA dictionary 2004: Summary of HLA-A, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR, and -DQ antigens. <i>Human Immunology</i> , 2005, 66, 170-210.	2.4	42
40	Race, Ethnicity and Ancestry in Unrelated Transplant Matching for the National Marrow Donor Program: A Comparison of Multiple Forms of Self-Identification with Genetics. <i>PLoS ONE</i> , 2015, 10, e0135960.	2.5	42
41	Genetic differentiation of Jewish populations. <i>Tissue Antigens</i> , 2010, 76, 442-458.	1.0	39
42	HLA class I haplotype diversity is consistent with selection for frequent existing haplotypes. <i>PLoS Computational Biology</i> , 2017, 13, e1005693.	3.2	38
43	Use of Cost-Effectiveness Analysis to Determine Inventory Size for a National Cord Blood Bank. <i>Medical Decision Making</i> , 2008, 28, 243-253.	2.4	36
44	World Marrow Donor Association guidelines for use of HLA nomenclature and its validation in the data exchange among hematopoietic stem cell donor registries and cord blood banks. <i>Bone Marrow Transplantation</i> , 2007, 39, 737-741.	2.4	35
45	HLA match likelihoods for Indian patients seeking unrelated donor transplantation grafts: a population-based study. <i>Lancet Haematology</i> , 2014, 1, e57-e63.	4.6	35
46	Hematopoietic stem cell donor registry strategies for assigning search determinants and matching relationships. <i>Bone Marrow Transplantation</i> , 2004, 33, 443-450.	2.4	34
47	Banking or Bankrupting: Strategies for Sustaining the Economic Future of Public Cord Blood Banks. <i>PLoS ONE</i> , 2015, 10, e0143440.	2.5	34
48	Fine-mapping of HLA associations with chronic lymphocytic leukemia in US populations. <i>Blood</i> , 2014, 124, 2657-2665.	1.4	33
49	Genetic editing of HLA expression in hematopoietic stem cells to broaden their human application. <i>Scientific Reports</i> , 2016, 6, 21757.	3.3	33
50	A comparative reference study for the validation of HLA matching algorithms in the search for allogeneic hematopoietic stem cell donors and cord blood units. <i>Hla</i> , 2016, 87, 439-448.	0.6	32
51	Significant variation between SNP-based HLA imputations in diverse populations: the last mile is the hardest. <i>Pharmacogenomics Journal</i> , 2018, 18, 367-376.	2.0	32
52	The HLA dictionary 2001: a summary of HLA-A, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR, and -DQ antigens. <i>Human Immunology</i> , 2001, 62, 826-849.	2.4	31
53	Identification by random forest method of HLA class I amino acid substitutions associated with lower survival at day 100 in unrelated donor hematopoietic cell transplantation. <i>Bone Marrow Transplantation</i> , 2012, 47, 217-226.	2.4	31
54	Histoimmunogenetics Markup Language 1.0: Reporting next generation sequencing-based HLA and KIR genotyping. <i>Human Immunology</i> , 2015, 76, 963-974.	2.4	30

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55	The HLA Dictionary 2001: a summary of HLA-A, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens. <i>Tissue Antigens</i> , 2001, 58, 109-140.	1.0	28
56	Minimum information for reporting next generation sequence genotyping (MIRING): Guidelines for reporting HLA and KIR genotyping via next generation sequencing. <i>Human Immunology</i> , 2015, 76, 954-962.	2.4	28
57	Haplotype associations of 90 rare alleles from the National Marrow Donor Program. <i>Tissue Antigens</i> , 2006, 67, 284-289.	1.0	27
58	HLA-A Disparities Illustrate Challenges for Ranking the Impact of HLA Mismatches on Bone Marrow Transplant Outcomes in the United States. <i>Biology of Blood and Marrow Transplantation</i> , 2009, 15, 971-981.	2.0	27
59	Comparative validation of computer programs for haplotype frequency estimation from donor registry data. <i>Tissue Antigens</i> , 2013, 82, 93-105.	1.0	26
60	Estimating KIR Haplotype Frequencies on a Cohort of 10,000 Individuals: A Comprehensive Study on Population Variations, Typing Resolutions, and Reference Haplotypes. <i>PLoS ONE</i> , 2016, 11, e0163973.	2.5	26
61	The HLA Dictionary 2001: a summary of HLA-A, -B, -C, -DRB1/3/4/5 and -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens. <i>International Journal of Immunogenetics</i> , 2001, 28, 565-596.	1.2	24
62	Large-scale DNA-based typing of HLA-A and HLA-B at low resolution is highly accurate specific and reliable. <i>Tissue Antigens</i> , 2000, 55, 352-358.	1.0	23
63	HLA Amino Acid Polymorphisms and Kidney Allograft Survival. <i>Transplantation</i> , 2017, 101, e170-e177.	1.0	23
64	Overview of registries, HLA typing and diversity, and search algorithms. <i>Tissue Antigens</i> , 2007, 69, 3-5.	1.0	22
65	Four-locus high-resolution HLA typing in a sample of Mexican Americans. <i>Tissue Antigens</i> , 2009, 74, 508-513.	1.0	22
66	Measuring Ambiguity in HLA Typing Methods. <i>PLoS ONE</i> , 2012, 7, e43585.	2.5	22
67	Charting improvements in US registry HLA typing ambiguity using a typing resolution score. <i>Human Immunology</i> , 2016, 77, 542-549.	2.4	21
68	World Marrow Donor Association framework for the implementation of HLA matching programs in hematopoietic stem cell donor registries and cord blood banks. <i>Bone Marrow Transplantation</i> , 2011, 46, 338-343.	2.4	20
69	Power Laws for Heavy-Tailed Distributions: Modeling Allele and Haplotype Diversity for the National Marrow Donor Program. <i>PLoS Computational Biology</i> , 2015, 11, e1004204.	3.2	20
70	The HLA dictionary 1999: a summary of HLA-A, -B, -C, -DRB1/3/4/5, -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR, and -DQ antigens. <i>Human Immunology</i> , 1999, 60, 1157-1181.	2.4	19
71	HLA polymorphism and risk of multiple myeloma. <i>Leukemia</i> , 2016, 30, 2260-2264.	7.2	19
72	Next generation sequencing characterizes the extent of HLA diversity in an Argentinian registry population. <i>Hla</i> , 2018, 91, 175-186.	0.6	19

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73	Consumer (dis-)interest in genetic ancestry testing: the roles of race, immigration, and ancestral certainty. <i>New Genetics and Society</i> , 2019, 38, 165-194.	1.2	19
74	The HLA Dictionary 2004: a summary of HLA-A, -B, -C, -DRB1/3/4/5 and -DQB1 alleles and their association with serologically defined HLA-A, -B, -C, -DR and -DQ antigens. <i>International Journal of Immunogenetics</i> , 2005, 32, 19-69.	1.8	18
75	Unrelated donor search prognostic score to support early HLA consultation and clinical decisions. <i>Bone Marrow Transplantation</i> , 2016, 51, 1476-1481.	2.4	18
76	Multiplicative fitness, rapid haplotype discovery, and fitness decay explain evolution of human MHC. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 14098-14104.	7.1	18
77	High-Resolution Match Rate of 7/8 and 9/10 or Better for the Be The Match Unrelated Donor Registry. <i>Biology of Blood and Marrow Transplantation</i> , 2016, 22, 759-763.	2.0	17
78	Use of a neural network to assign serologic specificities to HLA-A, -B and -DRB1 allelic products. <i>Tissue Antigens</i> , 2003, 62, 21-47.	1.0	16
79	16 th IHIW: Global analysis of registry HLA haplotypes from 20 Million individuals: Report from the IHIW Registry Diversity Group. <i>International Journal of Immunogenetics</i> , 2013, 40, 66-71.	1.8	16
80	High-resolution HLA A*14B*14DRB1 haplotype frequencies from the Ezer Mizion Bone Marrow Donor Registry in Israel. <i>Human Immunology</i> , 2016, 77, 1114-1119.	2.4	16
81	Collection and storage of HLA NGS genotyping data for the 17th International HLA and Immunogenetics Workshop. <i>Human Immunology</i> , 2018, 79, 77-86.	2.4	16
82	Maintaining updated DNA-based HLA assignments in the National Marrow Donor Program Bone Marrow Registry. <i>Reviews in Immunogenetics</i> , 2000, 2, 449-60.	0.7	16
83	Going back to the roots: effective utilisation of HLA typing information for bone marrow registries requires full knowledge of the DNA sequences of the oligonucleotide reagents used in the testing. <i>Tissue Antigens</i> , 2000, 56, 99-102.	1.0	15
84	Cord Blood Unit Access and Selection: 2010 and Beyond: Best Practices and Emerging Trends in Cord Blood Unit Selection. <i>Biology of Blood and Marrow Transplantation</i> , 2011, 17, S46-S51.	2.0	15
85	An update to the HLA Nomenclature Guidelines of the World Marrow Donor Association, 2012. <i>Bone Marrow Transplantation</i> , 2013, 48, 1387-1388.	2.4	14
86	Optimal Donor Selection for Hematopoietic Cell Transplantation Using Bayesian Machine Learning. <i>JCO Clinical Cancer Informatics</i> , 2021, 5, 494-507.	2.1	14
87	Stem cell donor HLA typing improves CPRA in kidney allocation. <i>American Journal of Transplantation</i> , 2021, 21, 138-147.	4.7	13
88	Information technology and the role of WMDA in promoting standards for international exchange of hematopoietic stem cell donors and products. <i>Bone Marrow Transplantation</i> , 2010, 45, 839-842.	2.4	12
89	Asymmetric linkage disequilibrium: Tools for assessing multiallelic LD. <i>Human Immunology</i> , 2016, 77, 288-294.	2.4	12
90	Improved accuracy of clinical HLA genotyping by next-generation DNA sequencing affects unrelated donor search results for hematopoietic stem cell transplantation. <i>Human Immunology</i> , 2018, 79, 848-854.	2.4	12

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91	Mapping molecular HLA typing data to UNOS antigen equivalents. <i>Human Immunology</i> , 2018, 79, 781-789.	2.4	12
92	GRIMM: GRaph IMputation and matching for HLA genotypes. <i>Bioinformatics</i> , 2019, 35, 3520-3523.	4.1	12
93	Chromosome Y-encoded antigens associate with acute graft-versus-host disease in sex-mismatched stem cell transplant. <i>Blood Advances</i> , 2018, 2, 2419-2429.	5.2	11
94	Reducing ethnic disparity in access to high-quality HLA-matched cord blood units for transplantation: analysis of the Canadian Blood Services' Cord Blood Bank inventory. <i>Transfusion</i> , 2019, 59, 2382-2388.	1.6	11
95	Efficient Sequencing, Assembly, and Annotation of Human KIR Haplotypes. <i>Frontiers in Immunology</i> , 2020, 11, 582927.	4.8	11
96	Demographic history and selection at HLA loci in Native Americans. <i>PLoS ONE</i> , 2020, 15, e0241282.	2.5	11
97	Investigating the Association of Genetic Admixture and Donor/Recipient Genetic Disparity with Transplant Outcomes. <i>Biology of Blood and Marrow Transplantation</i> , 2017, 23, 1029-1037.	2.0	10
98	Machine Learning Approach to Predicting Stem Cell Donor Availability. <i>Biology of Blood and Marrow Transplantation</i> , 2018, 24, 2425-2432.	2.0	10
99	HLA haplotype frequency estimation for heterogeneous populations using a graph-based imputation algorithm. <i>Human Immunology</i> , 2021, 82, 746-757.	2.4	10
100	16 th IHIW: Immunogenomic Data Management Methods. Report from the Immunogenomic Data Analysis Working Group (IDAWG). <i>International Journal of Immunogenetics</i> , 2013, 40, 46-53.	1.8	9
101	The association between HLA and non-Hodgkin lymphoma subtypes, among a transplant-indicated population. <i>Leukemia and Lymphoma</i> , 2019, 60, 2899-2908.	1.3	9
102	A Detailed View of KIR Haplotype Structures and Gene Families as Provided by a New Motif-Based Multiple Sequence Alignment. <i>Frontiers in Immunology</i> , 2020, 11, 585731.	4.8	9
103	The GL service: Web service to exchange GL string encoded HLA & KIR genotypes with complete and accurate allele and genotype ambiguity. <i>Human Immunology</i> , 2016, 77, 249-256.	2.4	8
104	A community standard XML message format for sequencing-based typing data. <i>Tissue Antigens</i> , 2007, 69, 69-71.	1.0	7
105	Identification of high-risk amino-acid substitutions in hematopoietic cell transplantation: a challenging task. <i>Bone Marrow Transplantation</i> , 2016, 51, 1342-1349.	2.4	7
106	Regarding "Recipients Receiving Better HLA-Matched Hematopoietic Cell Transplantation Grafts, Uncovered by a Novel HLA Typing Method, Have Superior Survival: A Retrospective Study" <i>Biology of Blood and Marrow Transplantation</i> , 2019, 25, e268-e269.	2.0	7
107	113-P. <i>Human Immunology</i> , 2006, 67, S127.	2.4	6
108	Prediction of HLA Genes from SNP Data and HLA Haplotype Frequencies. , 2012, , .		6

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109	East Meets West—Impact of Ethnicity on Donor Match Rates in the Ezer Mizion Bone Marrow Donor Registry. <i>Biology of Blood and Marrow Transplantation</i> , 2017, 23, 1381-1386.	2.0	6
110	Standard reference sequences for submission of <scp>HLA</scp> genotyping for the 18th International HLA and Immunogenetics Workshop. <i>Hla</i> , 2021, 97, 512-519.	0.6	6
111	Complementarity of Binding Motifs is a General Property of HLA-A and HLA-B Molecules and Does Not Seem to Effect HLA Haplotype Composition. <i>Frontiers in Immunology</i> , 2013, 4, 374.	4.8	5
112	Human leucocyte antigen (HLA)-A, -B, -C, -DRB1 and -DQB1 haplotype frequencies from 2491 cord blood units from Tamil speaking population from Tamil Nadu, India. <i>Molecular Biology Reports</i> , 2018, 45, 2821-2829.	2.3	5
113	Predicting HLA-DPB1 permissive probabilities through a DPB1 prediction service towards the optimization of HCT donor selection. <i>Human Immunology</i> , 2021, 82, 903-911.	2.4	5
114	Extensive haplotype diversity in African American mothers and their cord blood units. <i>Tissue Antigens</i> , 2013, 81, 28-34.	1.0	4
115	Modeling coverage gaps in haplotype frequencies via Bayesian inference to improve stem cell donor selection. <i>Immunogenetics</i> , 2018, 70, 279-292.	2.4	4
116	Assessment of HLA-B Genetic Variation with an HLA-B Leader Tool and Implications in Clinical Transplantation. <i>Blood Advances</i> , 2021, , .	5.2	4
117	Challenges for the standardized reporting of NGS HLA genotyping: Surveying gaps between clinical and research laboratories. <i>Human Immunology</i> , 2021, 82, 820-828.	2.4	4
118	Competing risks with missing covariates: effect of haplotypematch on hematopoietic cell transplant patients. <i>Lifetime Data Analysis</i> , 2013, 19, 19-32.	0.9	3
119	Donor Selection for Hematopoietic Stem Cell Transplant Using Cost-Sensitive SVM. , 2015, , .		3
120	High resolution HLA allele and haplotype frequencies for Arab donors in the Hadassah bone marrow donor registry. <i>Human Immunology</i> , 2019, 80, 823-827.	2.4	3
121	A new strategy for systematically classifying <scp>HLA</scp> alleles into serological specificities. <i>Hla</i> , 2022, 100, 193-231.	0.6	3
122	Diversity in exon 5 of HLA-Câ—04:01:01G is significant in anthropological studies. <i>Human Immunology</i> , 2016, 77, 426-428.	2.4	2
123	Single haplotype admixture models using large scale HLA genotype frequencies to reproduce human admixture. <i>Immunogenetics</i> , 2019, 71, 589-604.	2.4	2
124	Negative Impact of KIR-Ligand Mismatch on Transplant-Related Mortality (TRM) in Umbilical Cord Blood Transplant (UCBT) Recipients.. <i>Blood</i> , 2005, 106, 2041-2041.	1.4	2
125	Re-creation of the genetic composition of a founder population. <i>Human Genetics</i> , 2008, 124, 417-421.	3.8	1