

N P Mayor

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/6810135/publications.pdf>

Version: 2024-02-01

45
papers

841
citations

623734

14
h-index

501196

28
g-index

47
all docs

47
docs citations

47
times ranked

974
citing authors

#	ARTICLE	IF	CITATIONS
1	HLA Typing for the Next Generation. PLoS ONE, 2015, 10, e0127153.	2.5	125
2	HLA-DPB1 matching status has significant implications for recipients of unrelated donor stem cell transplants. Blood, 2006, 107, 1220-1226.	1.4	87
3	Recipients Receiving Better HLA-Matched Hematopoietic Cell Transplantation Grafts, Uncovered by a Novel HLA Typing Method, Have Superior Survival: A Retrospective Study. Biology of Blood and Marrow Transplantation, 2019, 25, 443-450.	2.0	84
4	Diverging effects of HLA-DPB1 matching status on outcome following unrelated donor transplantation depending on disease stage and the degree of matching for other HLA alleles. Leukemia, 2010, 24, 58-65.	7.2	83
5	Single molecule real-time DNA sequencing of HLA genes at ultra-high resolution from 126 International HLA and Immunogenetics Workshop cell lines. Hla, 2018, 91, 88-101.	0.6	59
6	Single Nucleotide Polymorphisms in the <i>NOD2/CARD15</i> Gene Are Associated With an Increased Risk of Relapse and Death for Patients With Acute Leukemia After Hematopoietic Stem-Cell Transplantation With Unrelated Donors. Journal of Clinical Oncology, 2007, 25, 4262-4269.	1.6	58
7	The degree of matching at HLA-DPB1 predicts for acute graft-versus-host disease and disease relapse following haematopoietic stem cell transplantation. Bone Marrow Transplantation, 2003, 31, 1001-1008.	2.4	49
8	Recipient/donor HLA and CMV matching in recipients of T-cell-depleted unrelated donor haematopoietic cell transplants. Bone Marrow Transplantation, 2017, 52, 717-725.	2.4	45
9	Identification of amyloid- β binding sites using an antisense peptide approach. NeuroReport, 2001, 12, 2561-2566.	1.2	33
10	Matching for 12 HLA Alleles Is Associated with a Significantly Superior Survival Due to a Lower Mortality in Recipients of Unrelated Donor Haematopoietic Cell Transplants for Early but Not Late Stage Leukaemia.. Blood, 2007, 110, 3056-3056.	1.4	30
11	Polymorphisms in the TNFA gene promoter region show evidence of strong linkage disequilibrium with HLA and are associated with delayed neutrophil engraftment in unrelated donor hematopoietic stem cell transplantation. Tissue Antigens, 2004, 63, 401-411.	1.0	21
12	Next-generation HLA typing of 382 International Histocompatibility Working Group reference B-lymphoblastoid cell lines: Report from the 17th International HLA and Immunogenetics Workshop. Human Immunology, 2019, 80, 449-460.	2.4	20
13	Impact of Previously Unrecognized HLA Mismatches Using Ultrahigh Resolution Typing in Unrelated Donor Hematopoietic Cell Transplantation. Journal of Clinical Oncology, 2021, 39, 2397-2409.	1.6	19
14	Particular HLA-DPB1 allele mismatches predict for worse overall survival in recipients of unrelated donor haematopoietic stem cell transplants. Biology of Blood and Marrow Transplantation, 2005, 11, 13.	2.0	15
15	<i>NOD2</i> Polymorphisms and Their Impact on Haematopoietic Stem Cell Transplant Outcome. Bone Marrow Research, 2012, 2012, 1-13.	1.7	11
16	Association of functional polymorphisms of the transforming growth factor B1 gene with survival and graft-versus-host disease after unrelated donor hematopoietic stem cell transplantation. Haematologica, 2010, 95, 276-283.	3.5	8
17	Presence of donor-encoded centromeric KIR B content increases the risk of infectious mortality in recipients of myeloablative, T-cell deplete, HLA-matched HCT to treat AML. Bone Marrow Transplantation, 2020, 55, 1975-1984.	2.4	8
18	Widespread non-coding polymorphism in <i>HLA</i> class II genes of International HLA and Immunogenetics Workshop cell lines. Hla, 2022, 99, 328-356.	0.6	7

#	ARTICLE	IF	CITATIONS
19	No impact of NOD2/CARD15 on outcome after SCT: a reply. Bone Marrow Transplantation, 2008, 42, 837-838.	2.4	6
20	Polymorphism in TGFBI is associated with worse non-relapse mortality and overall survival after stem cell transplantation with unrelated donors. Haematologica, 2016, 101, 382-390.	3.5	6
21	The novel <sc><i>HLA*03:04:01:47</i></sc> allele sequence identified using Pacific biosciences <sc>SMRT</sc> sequencing. Hla, 2020, 96, 525-526.	0.6	6
22	Extending the sequences of HLA class I alleles without full-length genomic coverage using single molecule real-time DNA sequencing. Hla, 2020, 95, 196-199.	0.6	6
23	Sequence of a novel HLA-A*0301 intronic variant (A*03010103). Tissue Antigens, 2005, 65, 107-109.	1.0	5
24	The novel <sc>HLA</sc>*44 allele, <sc>HLA</sc>*44:220</i>, identified by Single Molecule Real-time <sc>DNA</sc> sequencing in a British Caucasoid male. Tissue Antigens, 2015, 86, 61-63.	1.0	5
25	Single molecule real-time DNA sequencing of the full HLA-E gene for 212 reference cell lines. Hla, 2020, 95, 561-572.	0.6	5
26	KIR2DL1 allele sequence extensions and discovery of <i>2DL1*0010102</i> and <i>2DL1*0010103</i> alleles by DNA sequencing. Hla, 2018, 91, 546-547.	0.6	4
27	<i>HLA*DPB1*64:01N</i> and <i>DPB1*701:01</i> sequence extensions by single molecule real-time DNA sequencing. Hla, 2018, 92, 426-427.	0.6	4
28	Genomic sequence of the rare <sc>HLA</sc>*A*02:95</i> allele identified by sequence-based typing and cloning. Tissue Antigens, 2014, 84, 324-326.	1.0	3
29	Caspase-8 polymorphisms result in reduced Alemtuzumab-induced T-cell apoptosis and worse survival after transplantation. Bone Marrow Transplantation, 2015, 50, 237-243.	2.4	3
30	Characterization of the novel <sc>HLA</sc>*A*02:01:01:206</i> allele in a Northern European individual. Hla, 2022, 100, 59-60.	0.6	3
31	The novel <sc>HLA</sc>*DRB1</sc>*03:01:01:05</i> and <sc>DPB1</sc>*04:02:01:21</i> alleles identified in patients with acute leukemia. Hla, 2022, 99, 650-652.	0.6	3
32	Identification of the novel <sc>HLA</sc>*E</sc>*01</i>:<i>03</i>:<i>02</i>:<i>25</i> allele in an acute lymphoblastic leukemia patient. Hla, 2022, 100, 90-92.	0.6	3
33	A novel technique for NOD2/CARD15 genotyping using PCR-SSP. Journal of Immunological Methods, 2007, 327, 82-87.	1.4	2
34	Two novel alleles, <sc>HLA</sc>*A*32:01:01:09</i> and <sc>32:01:01:10</i>, identified by Pacific Bioscience's SMRT sequencing. Hla, 2018, 92, 409-411.	0.6	2
35	Identification of a new allele, <sc>HLA</sc>*DRB1*01:01:01:02</i> discovered using third-generation sequencing. Hla, 2019, 94, 458-459.	0.6	2
36	A genomic extension to the sequence of <sc>HLA</sc>*A*02:13</i>, identified using third-generation sequencing. Hla, 2019, 94, 437-438.	0.6	2

#	ARTICLE	IF	CITATIONS
37	A novel allele, <i>HLA*07:01:01:30</i> identified using third-generation sequencing. Hla, 2019, 94, 455-456.	0.6	2
38	The novel KIR2DL1 allele, <i>KIR2DL1*037</i>, defined in the cell line SPO010 (IHW9036). Hla, 2018, 91, 547-548.	0.6	1
39	Increased Disease Relapse in HLA-DPB1 Matched Recipients of Unrelated Donor Transplants Is Not Confined to Pairs Matched for the Other Five HLA loci.. Blood, 2004, 104, 829-829.	1.4	1
40	NOD2/CARD15 Genotype and HLA-DPB1 Matching Status Can Be Used To Predict Disease Relapse in Recipients of an Unrelated Donor Haematopoietic Stem Cell Transplant for Acute Leukaemia.. Blood, 2007, 110, 830-830.	1.4	1
41	High Resolution Allotyping of Four Alleles of HLA-DRB1*01 Group in Iranians Using Reverse-SSOPH Assay in Comparison with DNA Sequencing and PCR-SSP. Journal of Biological Sciences, 2008, 8, 392-397.	0.3	1
42	8: High recipient CD4+CD25hi regulatory T-cell level pre-transplant is associated with reduced overall survival after unrelated donor hematopoietic stem cell transplantation. Biology of Blood and Marrow Transplantation, 2007, 13, 5.	2.0	0
43	61: NOD2/CARD15 gene single nucleotide polymorphisms are associated with significant increases in mortality due to increases in disease relapse in recipients of an unrelated donor haematopoietic stem cell transplant for acute leukaemia. Biology of Blood and Marrow Transplantation, 2007, 13, 25.	2.0	0
44	Short template amplicon and multiplex megaprimer-enabled relay (STAMMER) sequencing, a simultaneous approach to higher throughput sequence-based typing of polymorphic genes. Immunogenetics, 2010, 62, 253-260.	2.4	0
45	HLA typing-A case-based approach to donor selection. Advances in Cell and Gene Therapy, 2018, 1, e16.	0.9	0