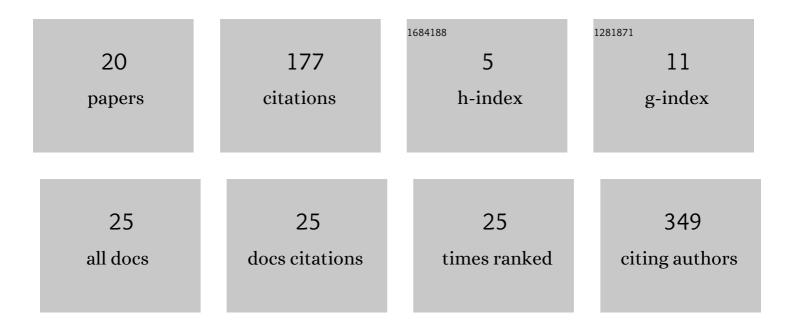
Junke Wang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/10197723/publications.pdf Version: 2024-02-01



LUNKE WANG

#	Article	IF	CITATIONS
1	Genome-Wide Non-HLA Mismatches Correlate with Overall Survival and Cause Specific Mortality after Unrelated Donor Allogeneic HCT. Transplantation and Cellular Therapy, 2021, 27, S61-S62.	1.2	Ο
2	Impact of Common Graft Versus Leukemia Minor Histocompatibility Antigens on the Outcomes of Allogeneic Hematopoietic Cell Transplantation (alloHCT). Transplantation and Cellular Therapy, 2021, 27, S156-S158.	1.2	0
3	Genome-Wide Association Analyses Identify Variants in IRF4 Associated With Acute Myeloid Leukemia and Myelodysplastic Syndrome Susceptibility. Frontiers in Genetics, 2021, 12, 554948.	2.3	8
4	Association of CXCR6 with COVID-19 severity: delineating the host genetic factors in transcriptomic regulation. Human Genetics, 2021, 140, 1313-1328.	3.8	33
5	Prognostic impact of pre-transplant chromosomal aberrations in peripheral blood of patients undergoing unrelated donor hematopoietic cell transplant for acute myeloid leukemia. Scientific Reports, 2021, 11, 15004.	3.3	4
6	Novel genetic variants associated with mortality after unrelated donor allogeneic hematopoietic cell transplantation. EClinicalMedicine, 2021, 40, 101093.	7.1	8
7	Pre-HCT mosaicism increases relapse risk and lowers survival in acute lymphoblastic leukemia patients post–unrelated HCT. Blood Advances, 2021, 5, 66-70.	5.2	6
8	Genome-wide association study identifies susceptibility loci for acute myeloid leukemia. Nature Communications, 2021, 12, 6233.	12.8	17
9	Prognostic Impact of Pre-Transplant Chromosomal Aberrations Detected By SNP-Array in Patients Undergoing Unrelated Donor Hematopoietic Cell Transplant for Acute Myeloid Leukemia. Biology of Blood and Marrow Transplantation, 2020, 26, S12-S13.	2.0	Ο
10	Meta-Analysis of Genome-Wide Association Studies of Acute Myeloid Leukemia (AML) Patients Identifies Variants Associated with Risk of 11q23/KMT2A-Translocated and Core-Binding Factor (CBF) AML and Suggests a Role for Transcription Elongation in Leukemogenesis. Blood, 2020, 136, 29-30.	1.4	0
11	Population Distribution of GvL and GvH Minor Histocompatibility Antigens. Blood, 2020, 136, 23-25.	1.4	0
12	Associations of Clinical Outcomes after Allogeneic Hematopoietic Cell Transplantation with Number of Predicted Class II Restricted mHA. Blood, 2020, 136, 2-2.	1.4	0
13	Pre-Transplant Clonal Mosaicism Is Associated with Increased Relapse and Lower Survival in Acute Lymphoblastic Leukemia Patients Undergoing Allogeneic Hematopoietic Cell Transplant. Blood, 2020, 136, 9-10.	1.4	Ο
14	Validation of genetic associations with acute GVHD and nonrelapse mortality in DISCOVeRY-BMT. Blood Advances, 2019, 3, 2337-2341.	5.2	8
15	Multiple functional variants in the IL1RL1 region are pretransplant markers for risk of GVHD and infection deaths. Blood Advances, 2019, 3, 2512-2524.	5.2	7
16	gwasurvivr: an R package for genome-wide survival analysis. Bioinformatics, 2019, 35, 1968-1970.	4.1	72
17	De Novo and Therapy-Related Acute Myeloid Leukemia and Myelodysplastic Syndrome: Similarities and Differences in SNP-Array Detected Chromosomal Aberrations in Pre-Transplant Blood Samples. Blood, 2019, 134, 1430-1430.	1.4	2
18	Genome Wide Interaction Analysis Identifies Expression Quantitative Trait Loci Associated with Reduced Survival after Reduced Intensity Conditioning HLA-Matched Unrelated Donor Allogeneic Hematopoietic Cell Transplant. Blood, 2019, 134, 4595-4595.	1.4	0

#	Article	IF	CITATIONS
19	Genome Wide Association Analyses Identify Pleiotropic Variants Associated with Acute Myeloid Leukemia (AML) and Myelodysplastic Syndrome (MDS) Susceptibility. Blood, 2018, 132, 1500-1500.	1.4	0
20	Multiple Functional Donor Polymorphisms in IL1RL1 region Associate with Death Due to GvHD or Infection after Unrelated Donor Allogeneic Hematopoietic Stem Cell Transplantation (HCT) for AML and MDS. Blood, 2018, 132, 312-312.	1.4	0