Yuelong Guo

List of Publications by Year in descending order

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687363 794594 20 950 13 19 citations h-index g-index papers 21 21 21 1811 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Ethnicity, sex, and age are determinants of red blood cell storage and stress hemolysis: results of the REDS-III RBC-Omics study. Blood Advances, 2017, 1, 1132-1141.	5. 2	164
2	Venetoclax responses of pediatric ALL xenografts reveal sensitivity of MLL-rearranged leukemia. Blood, 2016, 128, 1382-1395.	1.4	148
3	Donor sex, age and ethnicity impact stored red blood cell antioxidant metabolism through mechanisms in part explained by glucose 6-phosphate dehydrogenase levels and activity. Haematologica, 2021, 106, 1290-1302.	3.5	95
4	Genome-wide association study across European and African American ancestries identifies a SNP in DNMT3B contributing to nicotine dependence. Molecular Psychiatry, 2018, 23, 1911-1919.	7.9	80
5	Expanding the genetic architecture of nicotine dependence and its shared genetics with multiple traits. Nature Communications, 2020, 11, 5562.	12.8	80
6	Heterogeneity of blood processing and storage additives in different centers impacts stored red blood cell metabolism as much as storage time: lessons from REDSâ€IIIâ€"Omics. Transfusion, 2019, 59, 89-100.	1.6	71
7	Intradonor reproducibility and changes in hemolytic variables during red blood cell storage: results of recall phase of the REDSâ€III RBCâ€Omics study. Transfusion, 2019, 59, 79-88.	1.6	47
8	Frequent blood donations alter susceptibility of red blood cells to storage―and stress―induced hemolysis. Transfusion, 2019, 59, 67-78.	1.6	44
9	Blood, sweat, and tears: Red Blood Cellâ€Omics study objectives, design, and recruitment activities. Transfusion, 2019, 59, 46-56.	1.6	44
10	Clinical and genetic ancestry profile of a large multiâ€centre sickle cell disease cohort in Brazil. British Journal of Haematology, 2018, 182, 895-908.	2.5	38
11	OBI-3424, a Novel AKR1C3-Activated Prodrug, Exhibits Potent Efficacy against Preclinical Models of T-ALL. Clinical Cancer Research, 2019, 25, 4493-4503.	7.0	30
12	Development and evaluation of a transfusion medicine genome wide genotyping array. Transfusion, 2019, 59, 101-111.	1.6	30
13	Networks Underpinning Symbiosis Revealed Through Cross-Species eQTL Mapping. Genetics, 2017, 206, 2175-2184.	2.9	15
14	Preclinical evaluation of the combination of AZD1775 and irinotecan against selected pediatric solid tumors: A Pediatric Preclinical Testing Consortium report. Pediatric Blood and Cancer, 2020, 67, e28098.	1.5	13
15	Molecular phylogenetic analysis suggests paraphyly and early diversification of <i>Philadelphus</i> (<scp>H</scp> ydrangeaceae) in western North America: New insights into affinity with <i>Carpenteria</i> . Journal of Systematics and Evolution, 2013, 51, 545-563.	3.1	12
16	Genetic and behavioral modification of hemoglobin and iron status among firstâ€time and highâ€intensity blood donors. Transfusion, 2020, 60, 747-758.	1.6	9
17	Sex hormone intake in female blood donors: impact on haemolysis during cold storage and regulation of erythrocyte calcium influx by progesterone. Blood Transfusion, 2019, 17, 263-273.	0.4	9
18	Clinical and Genetic Predictors of Priapism in Sickle Cell Disease: Results from the Recipient Epidemiology and Donor Evaluation Study III Brazil Cohort Study. Journal of Sexual Medicine, 2019, 16, 1988-1999.	0.6	8

#	Article	IF	CITATIONS
19	Improved structural annotation of protein-coding genes in theMeloidogyne haplagenome using RNA-Seq. Worm, 2014, 3, e29158.	1.0	7
20	GWAS of Complete Blood Count (CBC) Measures in 13,403 Blood Donors in the Multi-Racial RBC-Omics Study Reveal Novel Genetic Loci in Minority Populations Which Provide Insights into the Pathways That May Connect Them to Disease. Blood, 2017, 130, 921-921.	1.4	0