Grier P Page

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8417568/publications.pdf Version: 2024-02-01

		394286	526166
30	1,316	19	27
papers	citations	h-index	g-index
33	33	33	2014
all docs	docs citations	times ranked	citing authors

CDIED D DACE

#	Article	IF	CITATIONS
1	Familial and Genetic Susceptibility to Major Neonatal Morbidities in Preterm Twins. Pediatrics, 2006, 117, 1901-1906.	1.0	298
2	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.	2.5	164
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.	1.7	95
4	Integrated Genomic Analyses in Bronchopulmonary Dysplasia. Journal of Pediatrics, 2015, 166, 531-537.e13.	0.9	93
5	Cis-Expression Quantitative Trait Loci Mapping Reveals Replicable Associations with Heroin Addiction in OPRM1. Biological Psychiatry, 2015, 78, 474-484.	0.7	64
6	Assessment of Genotype Imputation Performance Using 1000 Genomes in African American Studies. PLoS ONE, 2012, 7, e50610.	1.1	50
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.	0.8	47
8	Imputation across genotyping arrays for genome-wide association studies: assessment of bias and a correction strategy. Human Genetics, 2013, 132, 509-522.	1.8	44
9	Frequent blood donations alter susceptibility of red blood cells to storage―and stressâ€induced hemolysis. Transfusion, 2019, 59, 67-78.	0.8	44
10	Blood, sweat, and tears: Red Blood Cellâ€Omics study objectives, design, and recruitment activities. Transfusion, 2019, 59, 46-56.	0.8	44
11	Multiple-ancestry genome-wide association study identifies 27 loci associated with measures of hemolysis following blood storage. Journal of Clinical Investigation, 2021, 131, .	3.9	42
12	Family Environment, Neurodevelopmental Risk, and the Environmental Influences on Child Health Outcomes (ECHO) Initiative: Looking Back and Moving Forward. Frontiers in Psychiatry, 2020, 11, 547.	1.3	41
13	Large genome-wide association study identifies three novel risk variants for restless legs syndrome. Communications Biology, 2020, 3, 703.	2.0	40
14	Genes and environment in neonatal intraventricular hemorrhage. Seminars in Perinatology, 2015, 39, 592-603.	1.1	39
15	Genome-wide association study of sepsis in extremely premature infants. Archives of Disease in Childhood: Fetal and Neonatal Edition, 2017, 102, F439-F445.	1.4	32
16	Development and evaluation of a transfusion medicine genome wide genotyping array. Transfusion, 2019, 59, 101-111.	0.8	30
17	Blood donor obesity is associated with changes in red blood cell metabolism and susceptibility to hemolysis in cold storage and in response to osmotic and oxidative stress. Transfusion, 2021, 61, 435-448.	0.8	29
18	Donor genetic and nongenetic factors affecting red blood cell transfusion effectiveness. JCI Insight, 2022, 7, .	2.3	29

GRIER P PAGE

#	Article	IF	CITATIONS
19	Piloting and implementation of quality assessment and quality control procedures in RBCâ€Omics: a large multiâ€center study of red blood cell hemolysis during storage. Transfusion, 2019, 59, 57-66.	0.8	22
20	<i>KAT2B</i> polymorphism identified for drug abuse in African Americans with regulatory links to drug abuse pathways in human prefrontal cortex. Addiction Biology, 2016, 21, 1217-1232.	1.4	18
21	Surgical necrotizing enterocolitis in extremely premature neonates is associated with genetic variations in an intergenic region of chromosome 8. Pediatric Research, 2018, 83, 943-953.	1.1	17
22	Genetic variants associated with patent ductus arteriosus in extremely preterm infants. Journal of Perinatology, 2019, 39, 401-408.	0.9	16
23	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.	0.3	9
24	Genetic variation in dopamine neurotransmission and motor development of infants born extremely″owâ€birthweight. Developmental Medicine and Child Neurology, 2020, 62, 750-757.	1.1	3
25	Genetic predictors of severe intraventricular hemorrhage in extremely low-birthweight infants. Journal of Perinatology, 2021, 41, 286-294.	0.9	3
26	Sex-specific genetic modifiers identified susceptibility of cold stored red blood cells to osmotic hemolysis. BMC Genomics, 2022, 23, 227.	1.2	2
27	Evaluation of the Functional Effects of an African American Glucose-6-Phosphate Dehydrogenase (G6PD) Polymorphism (Val68Met) on RBC Hemolytic Propensity and Post-Transfusion Recovery in a Humanized Mouse Model. Blood, 2019, 134, 102-102.	0.6	1
28	In response. Transfusion, 2019, 59, 2750-2751.	0.8	0
29	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.	0.6	0
30	Demographic, Clinical, and Biochemical Predictors of Pica in a Large Cohort of Blood Donors. Blood, 2020, 136, 2-3.	0.6	0