Yanli Ji

List of Publications by Year in descending order

22 papers	189 citations	1163117 8 h-index	1125743 13 g-index
22 all docs	22 docs citations	22 times ranked	139 citing authors

#	Article	IF	Citations
1	Recommendation for validation and quality assurance of nonâ€invasive prenatal testing for foetal blood groups and implications for <scp>IVD</scp> risk classification according to <scp>EU</scp> regulations. Vox Sanguinis, 2022, 117, 157-165.	1.5	7
2	Incidence of <scp>antiâ€D</scp> alloimmunization in Dâ€negative individuals receiving Dâ€positive red blood cell transfusion: A systematic review and metaâ€analysis. Vox Sanguinis, 2022, 117, 633-640.	1.5	8
3	Psychological impact of the COVID â€19 pandemic on young professionals in blood banks and transfusion services: A global crossâ€sectional survey. Vox Sanguinis, 2022, , .	1.5	0
4	Secondary alloanti-D immunization post transfusion of â€œAsia typeâ€•DEL red blood cells. Transfusion and Apheresis Science, 2022, 61, 103458.	1.0	4
5	Serological screening and genetic analysis of RhCE variants in the Chinese Southern Han donors. Transfusion Medicine, 2021, 31, 271-276.	1.1	2
6	Hyporegenerative anemia in antiâ€Mâ€associated hemolytic disease of the fetus. Transfusion, 2021, 61, 1908-1915.	1.6	8
7	Molecular genetic analysis of Mi a â€positive hybrid glycophorins revealed two novel alleles of GP .Vw and multiple variant transcripts of GYPB existing in both the homozygous GP .Mur and wildâ€type GPB individuals. Transfusion, 2021, 61, 2477-2486.	1.6	0
8	Successful prenatal management of two foetuses affected by antibodies against GP.Mur with prenatal genotyping analysis and a literature review. Blood Transfusion, 2021, 19, 135-143.	0.4	1
9	Distribution of maternal red cell antibodies and the risk of severe alloimmune haemolytic disease of the foetus in a Chinese population: a cohort study on prenatal management. BMC Pregnancy and Childbirth, 2020, 20, 539.	2.4	13
10	Hemolytic disease of the fetus and newborn due to alloantiâ€M: three Chinese case reports and a review of the literature. Transfusion, 2019, 59, 385-395.	1.6	24
11	A variant RhAG protein encoded by theRHAG*572Aallele causes serological weak D expression while maintaining normal RhCE phenotypes. Transfusion, 2019, 59, 405-411.	1.6	7
12	Major applications and limitations of blood group genotyping in China. ISBT Science Series, 2018, 13, 365-370.	1.1	0
13	Genotyping analysis of MNS blood group GP(Bâ€Aâ€B) hybrid glycophorins in the Chinese Southern Han population using a highâ€resolution melting assay. Transfusion, 2018, 58, 1763-1771.	1.6	12
14	RHD genotype and zygosity analysis in the Chinese Southern Han D+, Dâ^' and D variant donors using the multiplex ligationâ€dependent probe amplification assay. Vox Sanguinis, 2017, 112, 660-670.	1.5	18
15	Validation of the multiplex ligationâ€dependent probe amplification assay and its application on the distribution study of the major alleles of 17 blood group systems in <scp>C</scp>hinese donors from <scp>G</scp>uangzhou. Transfusion, 2017, 57, 423-432.	1.6	8
16	The distribution of <scp>MNS</scp> hybrid glycophorins with Mur antigen expression in Chinese donors including identification of a novel <scp>GYP</scp>.Bun allele. Vox Sanguinis, 2016, 111, 308-314.	1.5	19
17	Identification of a novel frequentRHCE*ce308Tvariant allele in Chinese Dâ€“ individuals, resulting in a C+câ€“ phenotype. Transfusion, 2016, 56, 2314-2321.	1.6	2
18	Genotyping for Glycophorin GYP(B-A-B) Hybrid Genes Using a Single Nucleotide Polymorphism-Based Algorithm by Matrix-Assisted Laser Desorption/Ionisation, Time-of-Flight Mass Spectrometry. Molecular Biotechnology, 2016, 58, 665-671.	2.4	13

#	Article	IF	Citations
19	Red blood cell genotyping in China. ISBT Science Series, 2016, 11, 55-68.	1.1	6
20	Novel alleles at the <scp>K</scp>ell blood group locus that lead to <scp>K</scp>ell variant phenotype in the <scp>D</scp>utch population. Transfusion, 2015, 55, 413-421.	1.6	12
21	A novel 519_525dup mutation of <scp>KLF</scp>1 gene identified in a <scp>C</scp>hinese blood donor with <scp>L</scp>u(aâ€“bâ€“) phenotype. Transfusion, 2013, 53, 1619-1620.	1.6	7
22	The summary of <scp>FUT</scp>1 and <scp>FUT</scp>2 genotyping analysis in <scp>C</scp>hinese paraâ€<scp>B</scp>ombay individuals including additional nine probands from <scp>G</scp>uangzhou in <scp>C</scp>hina. Transfusion, 2013, 53, 3224-3229.	1.6	18