

# Robert L Flower

## List of Publications by Year in descending order

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Version: 2024-02-01

119  
papers

1,611  
citations

304743

22  
h-index

395702

33  
g-index

121  
all docs

121  
docs citations

121  
times ranked

1792  
citing authors

#	ARTICLE	IF	CITATIONS
1	In vitro antiviral activity of the anthraquinone chrysophanic acid against poliovirus. <i>Antiviral Research</i> , 2001, 49, 169-178.	4.1	134
2	Cytomegalovirus disease in immunocompetent adults. <i>Medical Journal of Australia</i> , 2014, 201, 578-580.	1.7	95
3	Evaluation of noninvasive prenatal RHD genotyping of the fetus. <i>Medical Journal of Australia</i> , 2009, 191, 21-25.	1.7	57
4	Evaluation of targeted exome sequencing for 28 protein-based blood group systems, including the homologous gene systems, for blood group genotyping. <i>Transfusion</i> , 2017, 57, 1078-1088.	1.6	54
5	A coarse-grained red blood cell membrane model to study stomatocyte-discocyte-echinocyte morphologies. <i>PLoS ONE</i> , 2019, 14, e0215447.	2.5	53
6	Canine Parvoviral Disease: Experimental Reproduction of the Enteric Form with a Parvovirus Isolated from a Case of Myocarditis. <i>Veterinary Pathology</i> , 1980, 17, 589-599.	1.7	51
7	Implications of Dengue Outbreaks for Blood Supply, Australia. <i>Emerging Infectious Diseases</i> , 2013, 19, 787-789.	4.3	51
8	Sixty Years of Antibodies to MNS System Hybrid Glycophorins: What Have We Learned?. <i>Transfusion Medicine Reviews</i> , 2011, 25, 111-124.	2.0	44
9	Hepatitis E Virus and Implications for Blood Supply Safety, Australia. <i>Emerging Infectious Diseases</i> , 2014, 20, 1940-1942.	4.3	34
10	Anti-D in pregnant women with the <i>i&gt;RHD&lt;/i&gt; (IVS3+1G&gt;A)-associated DEL phenotype. <i>Transfusion</i>, 2012, 52, 2016-2019.</i>	1.6	33
11	Investigation of red blood cell mechanical properties using AFM indentation and coarse-grained particle method. <i>BioMedical Engineering OnLine</i> , 2017, 16, 140.	2.7	31
12	IMMUNOHEMATOLOGY: Novel antibody screening cells, MUT+Mur kodeocytes, created by attaching peptides onto red blood cells. <i>Transfusion</i> , 2010, 50, 635-641.	1.6	30
13	Surface and Intracellular Interleukin-2 Receptor Expression on Various Resting and Activated Populations Involved in Cell-Mediated Immunity in Human Peripheral Blood. <i>Scandinavian Journal of Immunology</i> , 2000, 51, 67-72.	2.7	29
14	Hepatitis E virus <i>scp&gt;RNA&lt;/scp&gt;</i> in Australian blood donors: prevalence and risk assessment. <i>Vox Sanguinis</i> , 2017, 112, 614-621.	1.5	29
15	Approaches to Determination of a Full Profile of Blood Group Genotypes: Single Nucleotide Variant Mapping and Massively Parallel Sequencing. <i>Computational and Structural Biotechnology Journal</i> , 2014, 11, 147-151.	4.1	28
16	Targeted exome sequencing defines novel and rare variants in complex blood group serology cases for a red blood cell reference laboratory setting. <i>Transfusion</i> , 2018, 58, 284-293.	1.6	28
17	Hepatitis E virus RNA in Australian blood donations. <i>Transfusion</i> , 2016, 56, 3086-3093.	1.6	27
18	Procoagulant role of microparticles in routine storage of packed red blood cells: potential risk for prothrombotic post-transfusion complications. <i>Pathology</i> , 2017, 49, 62-69.	0.6	27

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19	The <i>RHD</i> (1227 <i>G</i> )&gt; <i>A</i> )</i>... <i>DEL</i> associated allele is the most prevalent <i>DEL</i> allele in Australian <i>D</i> blood donors with <i>C</i> + and/or <i>E</i> + phenotypes. <i>Transfusion</i> , 2014, 54, 2931-2940.	1.6	26
20	Soluble Mediators in Platelet Concentrates Modulate Dendritic Cell Inflammatory Responses in an Experimental Model of Transfusion. <i>Journal of Interferon and Cytokine Research</i> , 2015, 35, 821-830.	1.2	24
21	Antigenic differences between canine parvovirus and feline panleucopenia virus. <i>Veterinary Record</i> , 1980, 107, 254-256.	0.3	24
22	A Comparative Study of Assay Performance of Commercial Hepatitis E Virus Enzyme-Linked Immunosorbent Assay Kits in Australian Blood Donor Samples. <i>Journal of Blood Transfusion</i> , 2016, 2016, 1-6.	3.3	23
23	Assessment of In Vitro-Generated Platelet Microparticles Using a Modified Flow Cytometric Strategy. <i>Thrombosis Research</i> , 2001, 103, 47-55.	1.7	22
24	Noninvasive fetal <i>RHD</i> genotyping by microfluidics digital PCR using maternal plasma from two alloimmunized women with the variant <i>RHD</i> ( <i>IVS3+1G</i> &gt; <i>A</i> ) allele. <i>Prenatal Diagnosis</i> , 2013, 33, 1214-1216.	2.3	21
25	The distribution of <i>MNS</i> hybrid glycoporphins with Mur antigen expression in Chinese donors including identification of a novel <i>GYP</i> . <i>Bun</i> allele. <i>Vox Sanguinis</i> , 2016, 111, 308-314.	1.5	19
26	Diverse and novel <i>RHD</i> variants in Australian blood donors with a weak <i>D</i> phenotype: implication for transfusion management. <i>Vox Sanguinis</i> , 2017, 112, 279-287.	1.5	19
27	Seroprevalence of Antibodies to Ross River and Barmah Forest Viruses: Possible Implications for Blood Transfusion Safety After Extreme Weather Events. <i>EcoHealth</i> , 2015, 12, 347-353.	2.0	18
28	Genotyping confirms inheritance of the rare <i>At</i> ( <i>a</i> <sup>~</sup> ) type in a case of haemolytic disease of the newborn. <i>Journal of Pathology: Clinical Research</i> , 2016, 2, 53-55.	3.0	18
29	SPH-DEM approach to numerically simulate the deformation of three-dimensional RBCs in non-uniform capillaries. <i>BioMedical Engineering OnLine</i> , 2016, 15, 161.	2.7	18
30	Non-invasive fetal <i>RHD</i> genotyping for <i>RhD</i> negative women stratified into <i>RHD</i> gene deletion or variant groups: comparative accuracy using two blood collection tube types. <i>Pathology</i> , 2017, 49, 757-764.	0.6	18
31	Comprehensive blood group antigen profile predictions for Western Desert Indigenous Australians from whole exome sequence data. <i>Transfusion</i> , 2019, 59, 768-778.	1.6	18
32	Duffy blood group phenotype genotype correlations using high-resolution melting analysis <i>PCR</i> and microarray reveal complex cases including a new null <i>FY</i> * <i>A</i> allele: the role for sequencing in genotyping algorithms. <i>Vox Sanguinis</i> , 2015, 109, 296-303.	1.5	16
33	Non-invasive prenatal testing (NIPT) for fetal Kell, Duffy and <i>Rh</i> blood group antigen prediction in alloimmunised pregnant women: power of droplet digital PCR. <i>British Journal of Haematology</i> , 2020, 189, e90-e94.	2.5	16
34	<i>GYP</i> * <i>Kip</i> , a novel <i>GYP</i> ( <i>B</i> <i>A</i> <i>B</i> ) hybrid allele, encoding the <i>MNS</i> 48 ( <i>KIPP</i> ) antigen. <i>Transfusion</i> , 2016, 56, 539-541.	1.6	15
35	Hepatitis E virus seroepidemiology: a post-earthquake study among blood donors in Nepal. <i>BMC Infectious Diseases</i> , 2016, 16, 707.	2.9	15
36	Platelet concentrates modulate myeloid dendritic cell immune responses. <i>Platelets</i> , 2018, 29, 373-382.	2.3	15

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37	A novel <i>FY*</i> A allele with the 265T and 298A SNP's formerly associated exclusively with the <i>FY*</i> B allele and weak Fy <sup>b</sup> antigen expression: implication for genotyping interpretative algorithms. <i>Vox Sanguinis</i> , 2015, 108, 52-57.	1.5	14
38	An alloantibody in a homozygous <i>GYP*</i> Mur individual defines JENU (MNS49), a new high-frequency antigen on glycophorin B. <i>Transfusion</i> , 2017, 57, 716-717.	1.6	14
39	Noninvasive fetal RHD genotyping of RhD negative pregnant women for targeted anti-D therapy in Australia: A cost-effectiveness analysis. <i>Prenatal Diagnosis</i> , 2017, 37, 1245-1253.	2.3	14
40	Modelling of Red Blood Cell Morphological and Deformability Changes during In-Vitro Storage. <i>Applied Sciences (Switzerland)</i> , 2020, 10, 3209.	2.5	14
41	Coronary artery bypass grafting is associated with immunoparalysis of monocytes and dendritic cells. <i>Journal of Cellular and Molecular Medicine</i> , 2020, 24, 4791-4803.	3.6	14
42	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. <i>Molecular Biotechnology</i> , 2016, 58, 665-671.	2.4	13
43	SARA: a new low-frequency MNS antigen (MNS47) provides further evidence of the extreme diversity of the MNS blood group system. <i>Transfusion</i> , 2015, 55, 1451-1456.	1.6	12
44	Genotyping analysis of MNS blood group GP(B-A-B) hybrid glycoporphins in the Chinese Southern Han population using a high-resolution melting assay. <i>Transfusion</i> , 2018, 58, 1763-1771.	1.6	12
45	Non-invasive prenatal testing for management of haemolytic disease of the fetus and newborn induced by maternal alloimmunisation. <i>Transfusion and Apheresis Science</i> , 2020, 59, 102947.	1.0	12
46	Molecular typing for the Indian blood group associated 252G>C single nucleotide polymorphism in a selected cohort of Australian blood donors. <i>Blood Transfusion</i> , 2015, 13, 78-85.	0.4	12
47	Diffusion chamber method for in situ measurement of pathogen inactivation in groundwater. <i>Water Research</i> , 1998, 32, 1144-1150.	11.3	11
48	Frequency of <i>Mi</i> <sup>+</sup> <i>a</i> <sup>+</sup> (MNS7) and Classification of <i>Mi</i> <sup>+</sup> <i>a</i> <sup>+</sup> -Positive Hybrid Glycophorins in an Australian Blood Donor Population. <i>Transfusion Medicine and Hemotherapy</i> , 2020, 47, 279-287.	1.6	11
49	Targeted exome sequencing designed for blood group, platelet, and neutrophil antigen investigations: Proof-of-principle study for a customized single-test system. <i>Transfusion</i> , 2020, 60, 2108-2120.	1.6	11
50	Genetic factors associated with iron storage in Australian blood donors. <i>Blood Transfusion</i> , 2018, 16, 123-129.	0.4	11
51	Hepatitis E virus: do locally acquired infections in Australia necessitate laboratory testing in acute hepatitis patients with no overseas travel history?. <i>Pathology</i> , 2015, 47, 97-100.	0.6	10
52	A coupled SPH-DEM approach to model the interactions between multiple red blood cells in motion in capillaries. <i>International Journal of Mechanics and Materials in Design</i> , 2016, 12, 477-494.	3.0	10
53	Ross River virus in Australian blood donors: possible implications for blood transfusion safety. <i>Transfusion</i> , 2018, 58, 485-492.	1.6	10
54	Strategy for managing maternal variant RHD alleles in Rhesus D negative obstetric populations during fetal RHD genotyping. <i>Prenatal Diagnosis</i> , 2014, 34, 56-62.	2.3	9

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55	Mitigating the Risk of Transfusion-Transmitted Dengue in Australia. <i>Journal of Blood Transfusion</i> , 2016, 2016, 1-6.	3.3	9
56	A D+ blood donor with a novel RHD*DaCE(5aE6)âED gene variant exhibits the lowâ€frequency antigen RH23 (DW) characteristic of the partial DVa phenotype. <i>Transfusion</i> , 2016, 56, 2322-2330.	1.6	9
57	A proposed new lowâ€frequency antigen in the <sc>A</sc>ugustine blood group system associated with a severe case of hemolytic disease of the fetus and newborn. <i>Transfusion</i> , 2018, 58, 1320-1322.	1.6	9
58	Identification of genetic polymorphisms that predict responder/non-responder profiles to the RhD antigen. <i>Molecular Immunology</i> , 2015, 68, 628-633.	2.2	8
59	Immunomodulatory effect of cryopreserved platelets: altered BDCA3<sup>+</sup> dendritic cell maturation and activation in vitro. <i>Transfusion</i> , 2017, 57, 2878-2887.	1.6	8
60	Packed Red Blood Cell Transfusion Modulates Myeloid Dendritic Cell Activation and Inflammatory Response <i>In Vitro</i>. <i>Journal of Interferon and Cytokine Research</i> , 2018, 38, 111-121.	1.2	8
61	Severe hemolytic disease of the fetus and newborn due to alloâ€antiâ€D in a patient with a partial DEL phenotype arising from the variant allele described as <i>RHD*148+1T (RHD*01EL.31)</i>. <i>Transfusion</i> , 2018, 58, 2260-2264.	1.6	8
62	No evidence for widespread <i>Babesia microti</i> transmission in Australia. <i>Transfusion</i> , 2019, 59, 2368-2374.	1.6	8
63	HLA Antigens and the Response to Influenza A Virus. <i>Vox Sanguinis</i> , 1979, 37, 201-208.	1.5	7
64	A DEL phenotype attributed to <i>RHD</i> Exon 9 sequence deletion: slippedâ€strand mispairing and blood group polymorphisms. <i>Transfusion</i> , 2018, 58, 685-691.	1.6	7
65	Epidemic potential of Zika virus in Australia: implications for blood transfusion safety. <i>Transfusion</i> , 2019, 59, 648-658.	1.6	7
66	Glycophorins and the MNS blood group system: a narrative review. <i>Annals of Blood</i> , 0, 6, 39-39.	0.4	7
67	Epidemic Potential for Local Transmission of Zika Virus in 2015 and 2016 in Queensland, Australia. <i>PLOS Currents</i> , 2016, 8, .	1.4	7
68	A plasma ferritin is not always a serum ferritin. <i>Pathology</i> , 2015, 47, S89-S90.	0.6	6
69	Numerical Investigation of Motion and Deformation of a Single Red Blood Cell in a Stenosed Capillary. <i>International Journal of Computational Methods</i> , 2015, 12, 1540003.	1.3	6
70	Antiâ€D in a mother, hemizygous for the variant <i>RHD*DNB</i> gene, associated with hemolytic disease of the fetus and newborn. <i>Transfusion</i> , 2017, 57, 1938-1943.	1.6	6
71	Genetic Variants Within the Erythroid Transcription Factor, KLF1, and Reduction of the Expression of Lutheran and Other Blood Group Antigens: Review of the In(Lu) Phenotype. <i>Transfusion Medicine Reviews</i> , 2019, 33, 111-117.	2.0	6
72	RBCeq: A robust and scalable algorithm for accurate genetic blood typing. <i>EBioMedicine</i> , 2022, 76, 103759.	6.1	6

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73	Blood group genotype analysis of Australian reagent red blood cell donors across three genotyping platforms: consistent detection of 7Â-0% phenotype genotype nonconcordance. <i>ISBT Science Series</i> , 2014, 9, 309-314.	1.1	5
74	Cytomegalovirus in <sc>A</sc>ustralian blood donors: seroepidemiology and seronegative red blood cell component inventories. <i>Transfusion</i> , 2016, 56, 1616-1621.	1.6	5
75	Investigation of the variable In(Lu) phenotype caused by <i>KLF1</i> variants. <i>Transfusion</i> , 2018, 58, 2414-2420.	1.6	5
76	Mechanism of an action of an antibacterial murine lymphokine. <i>Nature</i> , 1975, 254, 459-460.	27.8	4
77	Identification of six new <i>RHCE</i> variant alleles in individuals of diverse racial origin. <i>Transfusion</i> , 2016, 56, 244-248.	1.6	4
78	Dâ€immunized blood donors who are female and who possess at least one HLAâ€DRB1*15 allele show a propensity for high serum RhIG production. <i>Transfusion</i> , 2018, 58, 1182-1188.	1.6	4
79	Emerging infectious disease outbreaks: estimating disease risk in Australian blood donors travelling overseas. <i>Vox Sanguinis</i> , 2018, 113, 21-30.	1.5	4
80	Estimation of mosquito-borne and sexual transmission of Zika virus in Australia: Risks to blood transfusion safety. <i>PLoS Neglected Tropical Diseases</i> , 2020, 14, e0008438.	3.0	4
81	Understanding occult hepatitis C infection. <i>Transfusion</i> , 2020, 60, 2144-2152.	1.6	4
82	The genomic landscape of blood groups in Indigenous Australians in remote communities. <i>Transfusion</i> , 2022, , .	1.6	4
83	Detection of emergent strains of <sc>W</sc>est <sc>N</sc>ile virus with a blood screening assay. <i>Transfusion</i> , 2016, 56, 1503-1507.	1.6	3
84	Genotyping by sequencing defines independent novel <i>RHD</i> variants for an antenatal patient and a blood donor. <i>Transfusion</i> , 2017, 57, 2281-2283.	1.6	3
85	Is Zika virus a potential threat to the Australian Blood Supply?. <i>Australian and New Zealand Journal of Public Health</i> , 2018, 42, 104-105.	1.8	3
86	Modeling the parvovirus B19 blood safety risk in Australia. <i>Transfusion</i> , 2019, 59, 295-302.	1.6	3
87	In Silico Analysis of Genetic Diversity of Human Hepatitis B Virus in Southeast Asia, Australia and New Zealand. <i>Viruses</i> , 2020, 12, 427.	3.3	3
88	Low Genetic Diversity of Hepatitis B Virus Surface Gene amongst Australian Blood Donors. <i>Viruses</i> , 2021, 13, 1275.	3.3	3
89	Hepatitis E virus infections in travellers: assessing the threat to the Australian blood supply. <i>Blood Transfusion</i> , 2017, 15, 191-198.	0.4	3
90	CHARACTERIZATION OF ENU-INDUCED MUTATIONS IN RED BLOOD CELL STRUCTURAL PROTEINS. <i>Computational and Structural Biotechnology Journal</i> , 2013, 6, e201303012.	4.1	2

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91	Incorporation of fluorescein conjugated function-spacer-lipid constructs into the red blood cell membrane facilitates detection of labeled cells for the duration of ex-vivo storage. <i>Journal of Immunological Methods</i> , 2016, 429, 66-70.	1.4	2
92	Whole-genome sequencing algorithm for blood-group typing. <i>Lancet Haematology</i> , 2018, 5, e233-e234.	4.6	2
93	Soluble mediators in packed red blood cells augment lipopolysaccharide-induced monocyte interleukin-1 $\beta$ production. <i>Vox Sanguinis</i> , 2020, 115, 562-569.	1.5	2
94	The interaction between Glycophorin A (GPA) and Band 3 in the formation of the Wright b (W <sup>r</sup> b <sup>s</sup> ) antigen. <i>Vox Sanguinis</i> , 2021, 116, 489-492.	1.5	2
95	Neonatal Outcomes From Arboviruses in the Perinatal Period: A State-of-the-Art Review. <i>Pediatrics</i> , 2021, 147, .	2.1	2
96	Molecular genotyping platforms for blood group antigen prediction. <i>Pathology</i> , 2014, 46, S87.	0.6	1
97	Next generation sequencing of an Australian family to identify the genetic basis of a rare blood group antigen. <i>Pathology</i> , 2014, 46, S87-S88.	0.6	1
98	Routine application of genotyping a step closer: direct PCR on plasma. <i>Annals of Blood</i> , 2021, 0, 2, 3-3.	0.4	1
99	GP.Mur red blood cells express variant form of s antigen (MNS4). <i>Pathology</i> , 2018, 50, S103.	0.6	1
100	Modified expression of the KEL2 (k) blood group antigen attributed to p.Leu196Val amino acid change three residues from the K/k antigen polymorphism site: implications for donor screening. <i>Transfusion</i> , 2019, 59, 1156-1158.	1.6	1
101	Inverse Relationship Between Lipopolysaccharide Concentration and Monocyte and Dendritic Cells Inflammatory Response. <i>Journal of Interferon and Cytokine Research</i> , 2020, 40, 349-356.	1.2	1
102	Computational modeling – an approach to the development of blood grouping reagents. <i>Expert Review of Hematology</i> , 2021, 14, 329-334.	2.2	1
103	Past and future epidemic potential of chikungunya virus in Australia. <i>PLoS Neglected Tropical Diseases</i> , 2021, 15, e0009963.	3.0	1
104	Using whole-genome sequencing to characterize clinically significant blood groups among healthy older Australians. <i>Blood Advances</i> , 2022, 6, 4593-4604.	5.2	1
105	Evaluation of testing strategies for reliable measurement of rates of subclinical mosquito-borne viral infections. <i>Pathology</i> , 2012, 44, S77.	0.6	0
106	Sero-Prevalence of antibodies to hepatitis A virus among Australian blood donors. <i>Pathology</i> , 2014, 46, S87.	0.6	0
107	Transfusion risk from emerging pathogens in the Asia-Pacific region. <i>ISBT Science Series</i> , 2016, 11, 143-148.	1.1	0
108	Hybrid glycophorins: Silent genetic variants complicate genetic testing. <i>Pathology</i> , 2016, 48, S98.	0.6	0

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109	Frequency of GP.Hut in association with â€™Mi(a)-positiveâ€™™ serology in Asian populations. Pathology, 2017, 49, S108.	0.6	0
110	Exposure to cryopreserved platelets mediates suppression of myeloid dendritic cell subset immune responses. Pathology, 2017, 49, S109.	0.6	0
111	Modulation of immune responses in patients following cardiac surgery. Pathology, 2017, 49, S109-S110.	0.6	0
112	Seroprevalence of antibodies to primate erythroparvovirus 1 among Australian blood donors. Pathology, 2017, 49, S115-S116.	0.6	0
113	KLF1 variants and the impact on the expression of red blood cell surface molecules in blood donors with the In(Lu) phenotype. Pathology, 2018, 50, S104.	0.6	0
114	Supernatants from packed red blood cells modulate inflammasome activation. Pathology, 2018, 50, S110-S111.	0.6	0
115	The role of nonâ€™invasive prenatal testing (NIPT) for fetal blood group typing in Australia. Australian and New Zealand Journal of Obstetrics and Gynaecology, 2021, , .	1.0	0
116	Increased circulating plasma mannose binding lectin associated serine protease (MASP) as a result of enu-induced mutation in MASP-1. Pathology, 2013, 45, S93.	0.6	0
117	Investigation of age, gender and MBL deficiency in relation to inflammatory markers in blood donors. Pathology, 2013, 45, S93.	0.6	0
118	An Improved Coarse-Grained Model to Accurately Predict Red Blood Cell Morphology and Deformability. , 2020, , 47-84.		0
119	A new highâ€™prevalence <sc>LW</sc> antigen detected by an antibody in an Indigenous Australian homozygous for <i> <sc>LW</sc> * <sc>A</sc> </i> c. <sc>309C</sc> &gt; <sc>A</sc> variant. Vox Sanguinis, 2022, , .	1.5	0