Jacques Chiaroni

List of Publications by Year
in descending order

147	3,746	159585 30	161849 54
papers	citations	h-index	g-index

181	181	181	5134
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	New insights into the Tyrolean Iceman's origin and phenotype as inferred by whole-genome sequencing. Nature Communications, 2012, 3, 698.	12.8	382
2	A major Y-chromosome haplogroup R1b Holocene era founder effect in Central and Western Europe. European Journal of Human Genetics, 2011, 19, 95-101.	2.8	224
3	Defining KIR and HLA Class I Genotypes at Highest Resolution via High-Throughput Sequencing. American Journal of Human Genetics, 2016, 99, 375-391.	6.2	156
4	Sequence family variant loss from the AZFc interval of the human Y chromosome, but not gene copy loss, is strongly associated with male infertility. Journal of Medical Genetics, 2004, 41, 814-825.	3.2	129
5	The phylogenetic and geographic structure of Y-chromosome haplogroup R1a. European Journal of Human Genetics, 2015, 23, 124-131.	2.8	122
6	Lower prevalence of antibodies neutralizing SARS-CoV-2 in group O French blood donors. Antiviral Research, 2020, 181, 104880.	4.1	121
7	Y chromosome diversity, human expansion, drift, and cultural evolution. Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 20174-20179.	7.1	115
8	Afghan Hindu Kush: Where Eurasian Sub-Continent Gene Flows Converge. PLoS ONE, 2013, 8, e76748.	2.5	96
9	Marseillevirus-Like Virus Recovered From Blood Donated by Asymptomatic Humans. Journal of Infectious Diseases, 2013, 208, 1042-1050.	4.0	93
10	HLA-DRB1 alleles and Jka immunization. Transfusion, 2005, 45, 956-959.	1.6	81
11	The <scp>HLA</scp>â€net G<scp>ENE[RATE</scp>] pipeline for effective <scp>HLA</scp> data analysis and its application to 145 population samples from Europe and neighbouring areas. Tissue Antigens, 2014, 83, 307-323.	1.0	79
12	Distinguishing the co-ancestries of haplogroup G Y-chromosomes in the populations of Europe and the Caucasus. European Journal of Human Genetics, 2012, 20, 1275-1282.	2.8	74
13	The emergence of Y-chromosome haplogroup J1e among Arabic-speaking populations. European Journal of Human Genetics, 2010, 18, 348-353.	2.8	71
14	Analysis of ABO discrepancies occurring in 35 French hospitals. Transfusion, 2004, 44, 860-864.	1.6	66
15	Genetic diversity on the Comoros Islands shows early seafaring as major determinant of human biocultural evolution in the Western Indian Ocean. European Journal of Human Genetics, 2011, 19, 89-94.	2.8	65
16	HLA-DRB1 polymorphism is associated with Kell immunisation. British Journal of Haematology, 2006, 132, 374-378.	2.5	64
17	Improving minority blood donation: anthropologic approach in a migrant community. Transfusion, 2007, 47, 402-409.	1.6	62
18	Red blood cell immunization in sickle cell disease: evidence of a large responder group and a low rate of anti-Rh linked to partial Rh phenotype. Haematologica, 2014, 99, e115-e117.	3.5	61

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19	Population structure of modern-day Italians reveals patterns of ancient and archaic ancestries in Southern Europe. Science Advances, 2019, 5, eaaw3492.	10.3	53
20	HLA-G haplotype structure shows good conservation between different populations and good correlation with high, normal and low soluble HLA-G expression. Human Immunology, 2013, 74, 203-206.	2.4	51
21	Prospective detection of chikungunya virus in blood donors, Caribbean 2014. Blood, 2014, 123, 3679-3681.	1.4	51
22	Molecular analysis of inactive and active RHD alleles in native Congolese cohorts. Transfusion, 2009, 49, 1353-1360.	1.6	48
23	Positive association of DRB1*04 and DRB1*15 alleles with Fy^a immunization in a Southern European population. Transfusion, 2009, 49, 2412-2417.	1.6	48
24	Epidemiology of Chikungunya Virus Outbreaks in Guadeloupe and Martinique, 2014: An Observational Study in Volunteer Blood Donors. PLoS Neglected Tropical Diseases, 2017, 11, e0005254.	3.0	44
25	Single PCR Multiplex SNaPshot Reaction for Detection of Eleven Blood Group Nucleotide Polymorphisms. Journal of Molecular Diagnostics, 2010, 12, 453-460.	2.8	43
26	Association of HLA-A and Non-Classical HLA Class I Alleles. PLoS ONE, 2016, 11, e0163570.	2.5	40
27	HLA-G UTR Haplotype Conservation in the Malian Population: Association with Soluble HLA-G. PLoS ONE, 2013, 8, e82517.	2.5	39
28	The coming of the Greeks to Provence and Corsica: Y-chromosome models of archaic Greek colonization of the western Mediterranean. BMC Evolutionary Biology, 2011, 11, 69.	3.2	37
29	Characterization of novel RHD alleles: relationship between phenotype, genotype, and trimeric architecture. Transfusion, 2012, 52, 2020-2029.	1.6	35
30	Weak D and DEL alleles detected by routine SNaPshot genotyping: identification of four novel RHD alleles. Transfusion, 2011, 51, 401-411.	1.6	34
31	Partial deletion in the JK locus causing a Jknull phenotype. Blood, 2002, 99, 1079-1081.	1.4	32
32	16th IHIW: Analysis of <scp>HLA</scp> Population Data, with updated results for 1996 to 2012 workshop data (<scp>AHPD</scp> project report). International Journal of Immunogenetics, 2013, 40, 21-30.	1.8	32
33	The Comoros Show the Earliest Austronesian Gene Flow into the Swahili Corridor. American Journal of Human Genetics, 2018, 102, 58-68.	6.2	32
34	Bronchial Epithelial Cells from Asthmatic Patients Display Less Functional HLA-G Isoform Expression. Frontiers in Immunology, 2017, 8, 6.	4.8	31
35	FCGR3A and FCGR2A Genotypes Differentially Impact Allograft Rejection and Patients' Survival After Lung Transplant. Frontiers in Immunology, 2019, 10, 1208.	4.8	29
36	HSFY genes and the P4 palindrome in the AZFb interval of the human Y chromosome are not required for spermatocyte maturation. Human Reproduction, 2012, 27, 615-624.	0.9	28

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37	Diversity of KIR, HLA Class I, and Their Interactions in Seven Populations of Sub-Saharan Africans. Journal of Immunology, 2019, 202, 2636-2647.	0.8	26
38	Identification of RHCE and KEL alleles in large cohorts of Afroâ€Caribbean and Comorian donors by multiplex SNaPshot and fragment assays: a transfusion support for sickle cell disease patients. British Journal of Haematology, 2011, 154, 260-270.	2.5	25
39	A genetic strategy to control expression of human blood group antigens in red blood cells generated in vitro. Transfusion, 2009, 49, 967-976.	1.6	24
40	Multiplex Lateral Flow Assay for Rapid Visual Blood Group Genotyping. Analytical Chemistry, 2018, 90, 7502-7509.	6.5	24
41	Distribution of killer-cell immunoglobulin-like receptor (KIR) in Comoros and Southeast France. Tissue Antigens, 2006, 67, 356-367.	1.0	23
42	Linkage disequilibrium between HLAâ€C*0104 and HLAâ€B*0103 alleles in Tswa Pygmies. Tissue Antigens, 2011, 77, 193-200.	1.0	23
43	HLAIb worldwide genetic diversity: New HLA-H alleles and haplotype structure description. Molecular Immunology, 2019, 112, 40-50.	2.2	23
44	A comprehensive survey of both <scp>RHD</scp> and <scp>RHCE</scp> allele frequencies in subâ€<scp>S</scp>aharan <scp>A</scp>frica. Transfusion, 2013, 53, 3009-3017.	1.6	22
45	Adsorption of autoantibodies in the presence of LISS to detect alloantibodies underlying warm autoantibodies. Transfusion, 2003, 43, 651-655.	1.6	20
46	Y-chromosome phylogeographic analysis of the Greek-Cypriot population reveals elements consistent with Neolithic and Bronze Age settlements. Investigative Genetics, 2016, 7, 1.	3.3	20
47	Validation of new HLAâ€F alleles assigned by nextâ€generation sequencing. Hla, 2019, 93, 131-132.	0.6	20
48	Correlation of annual precipitation with human Y-chromosome diversity and the emergence of Neolithic agricultural and pastoral economies in the Fertile Crescent. Antiquity, 2008, 82, 281-289.	1.0	19
49	Association between ABO haplotypes and the risk of venous thrombosis: impact on disease risk estimation. Blood, 2021, 137, 2394-2402.	1.4	19
50	Determination of the phylogenetic origins of the ÃrpÃ¡d Dynasty based on Y chromosome sequencing of BÃ©la the Third. European Journal of Human Genetics, 2021, 29, 164-172.	2.8	18
51	HLA-DRB1 and DQB1 polymorphisms in Southern France and genetic relationships with other Mediterranean populations. Human Immunology, 2000, 61, 930-936.	2.4	17
52	Genetic Characterization of the Population of Grande Comore Island (Njazidja) According to Major Blood Groups. Human Biology, 2004, 76, 527-541.	0.2	17
53	Analysis of hepatitis C virus strains circulating in Republic of the Congo. Journal of Medical Virology, 2010, 82, 562-567.	5.0	17
54	Evaluation of Next-Generation Sequencing and Crystal Digital PCR for Chimerism Monitoring of Post-Allogeneic Hematopoietic Stem Cell Transplantation. Transplantation and Cellular Therapy, 2021, 27, 89.e1-89.e10.	1.2	16

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55	Dombrock genotyping in a native Congolese cohort reveals two novel alleles. Transfusion, 2009, 49, 1661-1671.	1.6	15
56	Genetic data of 15 STR loci in five populations from Afghanistan. Forensic Science International: Genetics, 2012, 6, e44-e45.	3.1	15
57	A proposal for a rational transfusion strategy in patients of European and North African descent with weak D type 4.0 and 4.1 phenotypes. Blood Transfusion, 2019, 17, 89-90.	0.4	14
58	Duffy blood group genotyping in French Basques using polymerase chain reaction with alleleâ€specific primers (PCRâ€ASP). American Journal of Human Biology, 2004, 16, 78-81.	1.6	13
59	DNA-based typing of Kell, Kidd, MNS, Dombrock, Colton, and Yt blood group systems in the French Basques. American Journal of Human Biology, 2008, 20, 308-311.	1.6	13
60	Revisiting the Diego Blood Group System in Amerindians: Evidence for Gene-Culture Comigration. PLoS ONE, 2015, 10, e0132211.	2.5	13
61	A transcriptional signature associated with non-Hodgkin lymphoma in the blood of patients with Q fever. PLoS ONE, 2019, 14, e0217542.	2.5	13
62	Qualitative and quantitative comparison of cell-free DNA and cell-free fetal DNA isolation by four (semi-)automated extraction methods: impact in two clinical applications: chimerism quantification and noninvasive prenatal diagnosis. Journal of Translational Medicine, 2021, 19, 15.	4.4	13
63	Sub-30-nm hybrid lithography (electron beamâ•deep ultraviolet) and etch process for fully depleted metal oxide semiconductor transistors. Journal of Vacuum Science & Technology B, 2007, 25, 2030.	1.3	12
64	HLA-G Haplotypes Are Differentially Associated with Asthmatic Features. Frontiers in Immunology, 2018, 9, 278.	4.8	12
65	Development and validation of a genotyping kit for the eight principal human platelet alloantigen systems. Transfusion Clinique Et Biologique, 2000, 7, 51-62.	0.4	11
66	Use of the PSA enhancer core element to modulate the expression of prostate- and non-prostate-specific basal promoters in a lentiviral vector context. Cancer Gene Therapy, 2006, 13, 919-929.	4.6	11
67	Elimination of blood group antigens: hope and reality. British Journal of Haematology, 2011, 152, 392-400.	2.5	11
68	Identification of novel polymorphism restricted to the (C)ce^stype 1 haplotype avoids risk of transfusion deadlock in <scp>SCD</scp> patients. British Journal of Haematology, 2013, 160, 863-867.	2.5	11
69	Synonymous nucleotide polymorphisms influence <scp>D</scp>ombrock blood group protein expression in <scp>K</scp>562 cells. British Journal of Haematology, 2014, 164, 131-141.	2.5	11
70	RH diversity in Mali: characterization of a new haplotype RHD*DIVa/RHCE*ceTI(D2). Transfusion, 2015, 55, 1423-1431.	1.6	11
71	Responses of artificially reared cat fleas <scp>C</scp>tenocephalides felis felis (<scp>B</scp>ouchÃ©, 1835) to different mammalian bloods. Medical and Veterinary Entomology, 2015, 29, 171-177.	1.5	11
72	HLA-H: Transcriptional Activity and HLA-E Mobilization. Frontiers in Immunology, 2020, 10, 2986.	4.8	11

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73	Seroprevalence of hepatitis E virus among blood donors on Corsica, France, 2017. Eurosurveillance, 2020, 25, .	7.0	11
74	Neonatal Thrombocytopenia in HLAâ€ÐR, â€ÐQ, â€ÐPâ€Typed Mother due to Rare Antiâ€HPAâ€1b (PLA2) (Zwb) Fetomaternal Immunization. Vox Sanguinis, 1994, 67, 46-51.	1.5	10
75	ABO blood group, glycosyltransferase activity and risk of venous thromboembolism. Thrombosis Research, 2020, 193, 31-35.	1.7	10
76	Blood group typing in five Afghan populations in the North Hinduâ€Kush region: implications for blood transfusion practice. Transfusion Medicine, 2013, 23, 167-174.	1.1	9
77	Genome-wide analysis of Corsican population reveals a close affinity with Northern and Central Italy. Scientific Reports, 2019, 9, 13581.	3.3	9
78	Maternal HLA Ib Polymorphisms in Pregnancy Allo-Immunization. Frontiers in Immunology, 2021, 12, 657217.	4.8	9
79	Phase-shift mask for EUV lithography. , 2006, , .		8
80	How we evaluate panagglutinating sera. Transfusion, 2009, 49, 1540-1545.	1.6	8
81	High levels of molecular polymorphism at the KIR2DL4 locus in French and Congolese populations: Impact for anthropology and clinical studies. Human Immunology, 2009, 70, 953-959.	2.4	8
82	Molecular characterization of a new Dâ€•â€•haplotype in a Comorian man. Vox Sanguinis, 2012, 103, 352-355.	1.5	8
83	HLA-EâŽ01:03 Allele in Lung Transplant Recipients Correlates with Higher Chronic Lung Allograft Dysfunction Occurrence. Journal of Immunology Research, 2016, 2016, 1-8.	2.2	8
84	HLA-DRB1 frequencies of the Comorian population and their genetic affinities with Sub-Saharan African and Indian Oceanian populations. Annals of Human Biology, 2006, 33, 265-278.	1.0	7
85	The Araboâ€Islamic migrations in Madagascar: first genetic study of the GM system in three Malagasy populations. International Journal of Immunogenetics, 2012, 39, 161-169.	1.8	7
86	<scp>RHD</scp> zygosity assignments based on most probable genotype and hybrid <scp>Rhesus box</scp> detection in <scp>T</scp>unisia. Transfusion Medicine, 2012, 22, 362-366.	1.1	7
87	Genotyping of 28 blood group alleles in blood donors from Mali: Prediction of rare phenotypes. Transfusion and Apheresis Science, 2016, 54, 289-295.	1.0	7
88	Relevance and costs of RHD genotyping in women with a weak D phenotype. Transfusion Clinique Et Biologique, 2019, 26, 27-31.	0.4	7
89	Distribution of Rhesus Blood Group System in the French Basques: A Reappraisal Using the Allele-Specific Primers PCR Method. Human Heredity, 2004, 58, 69-72.	0.8	6
90	The use of the electronic (computer) cross-match. Vox Sanguinis, 2013, 104, 350-364.	1.5	6

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91	A simple genotyping procedure without <scp>DNA</scp> extraction to identify rare blood donors. Vox Sanguinis, 2015, 109, 173-180.	1.5	6
92	Prehistoric migrations through the Mediterranean basin shaped Corsican Y-chromosome diversity. PLoS ONE, 2018, 13, e0200641.	2.5	6
93	The radial expansion of the Diego blood group system polymorphisms in Asia: mark of co-migration with the Mongol conquests. European Journal of Human Genetics, 2019, 27, 125-132.	2.8	6
94	Paternal RHD zygosity determination in Tunisians: evaluation of three molecular tests. Blood Transfusion, 2015, 13, 59-65.	0.4	6
95	Groupes sanguins Ã©rythrocytaires. EMC - Hematologie, 2005, 2, 53-112.	0.1	5
96	Heterogeneity of alleles encoding highâ€•and lowâ€prevalence red blood cell antigens across <scp>A</scp>frica: useful data to facilitate transfusion in <scp>A</scp>frican patients. British Journal of Haematology, 2013, 163, 528-536.	2.5	5
97	Unreliable patient identification warrants ABO typing at admission to check existing records before transfusion. Transfusion Clinique Et Biologique, 2015, 22, 66-70.	0.4	5
98	Short duplication within the <scp>RHCE</scp> gene associated with an in cis deleted <scp>RHD</scp> causing a <scp>R</scp>h_{null} amorph phenotype in an immunized pregnant woman with antiâ€<scp>R</scp>h29. Transfusion, 2015, 55, 1407-1410.	1.6	5
99	Comparing two blood culture systems for the detection of bacterial contamination in platelet concentrates. Transfusion, 2018, 58, 2604-2610.	1.6	5
100	Validation of new <scp>HLAâ€J</scp> alleles assigned by next generation sequencing. Hla, 2021, 98, 173-175.	0.6	5
101	Blood groups of Neandertals and Denisova decrypted. PLoS ONE, 2021, 16, e0254175.	2.5	5
102	HLA-H*02:07 Is a Membrane-Bound Ligand of Denisovan Origin That Protects against Lysis by Activated Immune Effectors. Journal of Immunology, 2022, 208, 49-53.	0.8	5
103	Analysis of a large single institution cohort of related donors fails to detect a relation between SDF1/CXCR4 or VCAM/VLA4 genetic polymorphisms and the level of hematopoietic progenitor cell mobilization in response to G-CSF. PLoS ONE, 2020, 15, e0228878.	2.5	4
104	Complete genetic sequence of 15 novel HLAâ€H alleles. Hla, 2020, 96, 133-135.	0.6	4
105	Identification of 11 novel <scp>HLAâ€A</scp>, â€B, â€C, â€<scp>DRB1,</scp> and â€<scp>DQB1</scp> alleles in the 1000 Genomes Project panel. Hla, 2021, 97, 246-248.	0.6	4
106	Ã‰tude comparative de diffÃ©rents procÃ©dÃ©s de filtration : application Ã la recherche des anticorps irrÃ©guliers anti-Ã©rythrocytaires (RAI). Revue Francaise Des Laboratoires, 2000, 2000, 33-36.	0.0	3
107	Risque immuno-hÃ©molytique des transfusions sanguines et analyses d'immuno-hÃ©matologieÃ©rythrocytaire. Revue Francaise Des Laboratoires, 2003, 2003, 45-51.	0.0	3
108	RHCE*cE734C allele encodes an altered c antigen and a suppressed E antigen not detected with standard reagents. Transfusion, 2013, 53, 955-961.	1.6	3

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109	Subtle adjustments of the glucose-6-phosphate dehydrogenase (G6PD) mutation database and reference sequence. Blood Cells, Molecules, and Diseases, 2014, 52, 55-56.	1.4	3
110	New silent and weak D alleles: molecular characterization and associated antigen density. Transfusion, 2016, 56, 2154-2155.	1.6	3
111	CORS (CROM20): A new highâ€prevalence antigen in the Cromer blood group system. Transfusion, 2020, 60, E40-E42.	1.6	3
112	In vitro detection of bacterial contamination in platelet concentrates by matrix-assisted laser desorption/ionization time-of-flight mass spectrometry: a preliminary study. Journal of Medical Microbiology, 2017, 66, 1523-1530.	1.8	3
113	Droplet Digital PCR: A New Technology for Detection and Quantification of Chimerism After Allogenic Hematopoietic Stem Cell Transplantation. Biomedical Journal of Scientific & Technical Research, 2019, 13, .	0.1	3
114	The above letter was sent to Dr Ferrera et al.: Drs Ferrera, Chiaroni, Legrand, and Dettori. Transfusion, 2005, 45, 455-455.	1.6	2
115	Off-on polyadenylation strategy as a supplemental mechanism for silencing toxic transgene expression during lentiviral vector production. BioTechniques, 2014, 56, 311-2, 314-8.	1.8	2
116	Sub-Saharan red cell antigen phenotypes and glucose-6-phosphate dehydrogenase deficiency variants in French Guiana. Malaria Journal, 2016, 15, 310.	2.3	2
117	Genes flow by the channels of culture: the genetic imprint of matrilocality in Ngazidja, Comoros Islands. European Journal of Human Genetics, 2018, 26, 1222-1226.	2.8	2
118	Rapid identification of microorganisms from platelet concentrates by matrixâ€assisted laser desorption ionization timeâ€ofâ€flight mass spectrometry after shortâ€term incubation on liquid medium. Transfusion, 2018, 58, 766-773.	1.6	2
119	DO/ART4 gene sequencing in subâ€Saharan cohorts and African migrants: useful data describing the diversity and spreading of rare variants. Transfusion, 2019, 59, 3755-3766.	1.6	2
120	Presence of SARS-CoV-2 in a Cornea Transplant. Pathogens, 2021, 10, 934.	2.8	2
121	Blood group typing from whole-genome sequencing data. PLoS ONE, 2020, 15, e0242168.	2.5	2
122	Impact of societal and legal context on the blood supply of Africanâ€ancestry populations in Western countries: A review of practices and the French example. Vox Sanguinis, 2022, 117, 1137-1144.	1.5	2
123	Genetic polymorphism (ABO, Rh, Kell) in Kabul (Afghanistan). International Journal of Anthropology, 2001, 16, 275-280.	0.1	1
124	ABO discrepancy by usurpation of identity. Transfusion, 2005, 45, 454-454.	1.6	1
125	DÃ©terminants gÃ©nÃ©tiques deÂlaÂrÃ©ponse auÂclopidogrel. Hematologie, 2009, 15, 045-071.	0.0	1
126	La sÃ©curitÃ© immuno-hÃ©matologique desÂreceveurs. Hematologie, 2010, 16, 156-161.	0.0	1

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127	Common subclinical hypothyroidism during Whippleâ€™s disease. BMC Infectious Diseases, 2014, 14, 370.	2.9	1
128	Sequencing of the ART4 gene in subâ€Saharan cohorts reveals ethnic differences and two new DO alleles: DO*Bâ€Ile5Thr and DO*Bâ€Trp266Arg. Transfusion, 2015, 55, 2376-2383.	1.6	1
129	New KEL*01M and KEL*02M alleles: structural modeling to assess the impact of amino acid changes. Transfusion, 2016, 56, 1223-1229.	1.6	1
130	Genetic polymorphisms with erythrocyte traits in malaria endemic areas of Mali. PLoS ONE, 2019, 14, e0209966.	2.5	1
131	Maternal HLA-G*01:01:01:04 protects from anti-HLA-class II immunization in pregnant women. Human Immunology, 2019, 80, 120-125.	2.4	1
132	Lâ€™Ã©tablissement franÃ§ais du sang Alpes-MÃ©diterranÃ©e, une structure de santÃ© doublÃ©e dâ€™un observatoire de la diversitÃ© populationnelle marseillaise. Cahiers QuÃ©bÃ©cois De DÃ©mographie, 0, 36, 85-110.	0.5	1
133	Identification of Frailty in a Population of Former Immigrant Workers in the South of France. Journal of Nutrition, Health and Aging, 2021, 25, 1226-1228.	3.3	1
134	Ã‰valuation externe de la qualitÃ©. Transfusion Clinique Et Biologique, 2001, 8, 475-477.	0.4	0
135	aide Ã la dÃ©cision en immunohÃ©matologie : Ã©preuve directe de comptabilitÃ© au laboratoire (edc). Transfusion Clinique Et Biologique, 2001, 8, 481-484.	0.4	0
136	Integration issues in step and repeat UV nanoimprint lithography. , 2008, , .		0
137	LETTERS TO THE EDITOR: Comparison of three lowâ€ionicâ€strength solutions for routine pretransfusion testing: antibody screening/identification, crossâ€matching, immune antiâ€ABO detection, and direct antiglobulin tests. Transfusion, 2009, 49, 2772-2773.	1.6	0
138	Transfusion sanguine: dÃ©bats d'actualitÃ© 2010. Hematologie, 2010, 16, 29-46.	0.0	0
139	Correction for Chiaroni et al., Y chromosome diversity, human expansion, drift, and cultural evolution. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 13556-13556.	7.1	0
140	Blood transfusion: current debates. Hematologie, 2011, 17, 71-87.	0.0	0
141	First description of a Dâ€CEâ€D hybrid gene on a weak D Type 2 molecular background. Transfusion, 2017, 57, 1248-1253.	1.6	0
142	Transfusion practices in geriatric short stay unit before and after the French national health authority guidelines of 2014. Psychologie & Neuropsychiatrie Du Vieillissement, 2018, 16, 367-375.	0.2	0
143	Identification and characterization of three novel RHCE*ce variant alleles affecting Rhc (RH4) reactivity. Transfusion, 2019, 59, 2754-2755.	1.6	0
144	The c.939G>A synonymous polymorphism in RHCE can be encountered on different molecular backgrounds. Transfusion, 2019, 59, 2160-2161.	1.6	0

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145	Giving blood: providing treatment, providing knowledge. European Journal of Public Health, 2019, 29, .	0.3	0
146	RHCE*01 (<scp>c.499A>G</scp>,<scp>p.Met167Val</scp>)allele: Weak RhE expression which does not require the Eâ€specific proline 226. Transfusion, 2021, 61, E18-E20.	1.6	0
147	Five-Years Review of RHCE Alleles Detected after Weak and/or Discrepant C Results in Southern France. Genes, 2022, 13, 1058.	2.4	0