Erick C Castelli

List of Publications by Year in descending order

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186209 189801 2,977 100 28 50 citations h-index g-index papers 113 113 113 2148 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Implications of the polymorphism of HLA-G on its function, regulation, evolution and disease association. Cellular and Molecular Life Sciences, 2011, 68, 369-395.	2.4	302
2	The genetic structure of $3\hat{a}\in^2$ untranslated region of the HLA-G gene: polymorphisms and haplotypes. Genes and Immunity, 2010, 11, 134-141.	2.2	182
3	Transcriptional and Posttranscriptional Regulations of the <i>HLA-G</i> Gene. Journal of Immunology Research, 2014, 2014, 1-15.	0.9	156
4	A Comprehensive Study of Polymorphic Sites along the HLA-G Gene: Implication for Gene Regulation and Evolution. Molecular Biology and Evolution, 2011, 28, 3069-3086.	3 . 5	142
5	In silico analysis of microRNAS targeting the HLA-G 3′ untranslated region alleles and haplotypes. Human Immunology, 2009, 70, 1020-1025.	1.2	139
6	Polymorphic Sites at the 3' Untranslated Region of the HLA-G Gene Are Associated with Differential hla-g Soluble Levels in the Brazilian and French Population. PLoS ONE, 2013, 8, e71742.	1.1	139
7	Insights into HLA-G Genetics Provided by Worldwide Haplotype Diversity. Frontiers in Immunology, 2014, 5, 476.	2.2	119
8	Emerging role of PTEN loss in evasion of the immune response to tumours. British Journal of Cancer, 2020, 122, 1732-1743.	2.9	95
9	Worldwide genetic variation at the 3′ untranslated region of the HLA-G gene: balancing selection influencing genetic diversity. Genes and Immunity, 2014, 15, 95-106.	2.2	69
10	microRNAs targeting the immunomodulatory HLA-G gene: A new survey searching for microRNAs with potential to regulate HLA-G. Molecular Immunology, 2015, 65, 230-241.	1.0	61
11	Neither self-reported ethnicity nor declared family origin are reliable indicators of genomic ancestry. Genetica, 2016, 144, 259-265.	0.5	58
12	The Role of HLA-G Molecule and HLA-G Gene Polymorphisms in Tumors, Viral Hepatitis, and Parasitic Diseases. Frontiers in Immunology, 2015, 6, 9.	2.2	55
13	Hla-mapper: An application to optimize the mapping of HLA sequences produced by massively parallel sequencing procedures. Human Immunology, 2018, 79, 678-684.	1.2	51
14	HLA-G alleles and HLA-G 14 bp polymorphisms in a Brazilian population. Tissue Antigens, 2007, 70, 62-68.	1.0	50
15	HLA-G polymorphisms in women with squamous intraepithelial lesions harboring human papillomavirus. Modern Pathology, 2009, 22, 1075-1082.	2.9	48
16	HLA-G 3â€ ² UTR polymorphisms in high grade and invasive cervico-vaginal cancer. Human Immunology, 2013, 74, 452-458.	1.2	47
17	<i>HLAâ€G </i> polymorphism and transitional cell carcinoma of the bladder in a Brazilian population. Tissue Antigens, 2008, 72, 149-157.	1.0	46
18	HLA-G $3\hat{a}\in^2$ UTR-2 haplotype is associated with Human African trypanosomiasis susceptibility. Infection, Genetics and Evolution, 2013, 17, 1-7.	1.0	42

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19	Haplotypes of the HLA-G 3' Untranslated Region Respond to Endogenous Factors of HLA-G+ and HLA-G-Cell Lines Differentially. PLoS ONE, 2017, 12, e0169032.	1.1	39
20	Association of HLA-G 3′UTR polymorphisms with response to malaria infection: A first insight. Infection, Genetics and Evolution, 2013, 16, 263-269.	1.0	35
21	Genetic association between HLA-G 14-bp polymorphism and diseases: A systematic review and meta-analysis. Human Immunology, 2018, 79, 724-735.	1.2	35
22	MHC Variants Associated With Symptomatic Versus Asymptomatic SARS-CoV-2 Infection in Highly Exposed Individuals. Frontiers in Immunology, 2021, 12, 742881.	2.2	35
23	Whole-genome sequencing of 1,171 elderly admixed individuals from Brazil. Nature Communications, 2022, 13, 1004.	5.8	35
24	Frequency of insertion/deletion polymorphism in exon 8 of <i>HLAâ€G</i> and kidney allograft outcome. Tissue Antigens, 2008, 71, 35-41.	1.0	34
25	HLA-G 14-bp polymorphism at exon 8 in Amerindian populations from the Brazilian Amazon. Tissue Antigens, 2007, 69, 255-260.	1.0	33
26	Worldwide <i><scp>HLA</scp>â€E</i> nucleotide and haplotype variability reveals a conserved gene for coding and 3′ untranslated regions. Tissue Antigens, 2014, 83, 82-93.	1.0	33
27	HLA-E coding and 3′ untranslated region variability determined by next-generation sequencing in two West-African population samples. Human Immunology, 2015, 76, 945-953.	1.2	33
28	Ancestry informative markers and selected single nucleotide polymorphisms in immunoregulatory genes on preterm labor and preterm premature rupture of membranes: a case control study. BMC Pregnancy and Childbirth, 2016, 16, 30.	0.9	33
29	HLA-G 3′ Untranslated Region Polymorphisms Are Associated with Systemic Lupus Erythematosus in 2 Brazilian Populations. Journal of Rheumatology, 2013, 40, 1104-1113.	1.0	32
30	Haplotype frequencies based on eight polymorphic sites at the $3\hat{a} \in \mathbb{C}^2$ untranslated region of the $\langle i \rangle$ HLA $\hat{a} \in \mathbb{C} \langle i \rangle$ gene in individuals from two different geographical regions of Brazil. Tissue Antigens, 2012, 79, 272-278.	1.0	31
31	HLA-G variability and haplotypes detected by massively parallel sequencing procedures in the geographicaly distinct population samples of Brazil and Cyprus. Molecular Immunology, 2017, 83, 115-126.	1.0	29
32	Association of <i>HLAâ€G</i> 3′ untranslated region polymorphisms with antibody response against <i>Plasmodium falciparum</i> antigens: preliminary results. Tissue Antigens, 2013, 82, 53-58.	1.0	28
33	HLA-F coding and regulatory segments variability determined by massively parallel sequencing procedures in a Brazilian population sample. Human Immunology, 2016, 77, 841-853.	1.2	28
34	DC-SIGN (CD209) gene promoter polymorphisms in a Brazilian population and their association with human T-cell lymphotropic virus type 1 infection. Journal of General Virology, 2009, 90, 927-934.	1.3	25
35	Human leukocyte antigen–G 3′ untranslated region polymorphisms are associated with better kidney allograft acceptance. Human Immunology, 2012, 73, 52-59.	1.2	24
36	Balancing immunity and tolerance: genetic footprint of natural selection in the transcriptional regulatory region of HLA-G. Genes and Immunity, 2015, 16, 57-70.	2.2	24

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37	Nonâ€classical <i>HLAâ€E</i> gene variability in Brazilians: a nearly invariable locus surrounded by the most variable genes in the human genome. Tissue Antigens, 2012, 79, 15-24.	1.0	22
38	Insights on the HLA-G Evolutionary History Provided by a Nearby Alu Insertion. Molecular Biology and Evolution, 2013, 30, 2423-2434.	3.5	22
39	Evaluation of computational methods for the reconstruction of HLA haplotypes. Tissue Antigens, 2010, 76, 459-466.	1.0	21
40	A single nucleotide deletion at the C1 inhibitor gene as the cause of hereditary angioedema: insights from a Brazilian family. Allergy: European Journal of Allergy and Clinical Immunology, 2011, 66, 1384-1390.	2.7	20
41	Association of HLA-G 3′ untranslated region variants with type 1 diabetes mellitus. Human Immunology, 2016, 77, 358-364.	1.2	20
42	HLA-F displays highly divergent and frequent haplotype lineages associated with different mRNA expression levels. Human Immunology, 2019, 80, 112-119.	1.2	20
43	Haplotypes from the SLC45A2 gene are associated with the presence of freckles and eye, hair and skin pigmentation in Brazil. Legal Medicine, 2017, 25, 43-51.	0.6	19
44	Interleukin-18 and interferon-gamma polymorphisms in Brazilian human immunodeficiency virus-1-infected patients presenting with lipodystrophy syndrome. Tissue Antigens, 2010, 76, 126-30.	1.0	18
45	Immunogenetics of prostate cancer and benign hyperplasia $\hat{a} \in \text{``the potential use of an HLA} \in G$ variant as a tag SNP for prostate cancer risk. Hla, 2016, 87, 79-88.	0.4	18
46	Genetic diversity of the HLA-G coding region in Amerindian populations from the Brazilian Amazon: a possible role of natural selection. Genes and Immunity, 2013, 14, 518-526.	2.2	17
47	Absence of the HLA-G*0105N allele in Amerindian populations from the Brazilian Amazon Region: a possible role of natural selection. Tissue Antigens, 2007, 70, 330-334.	1.0	16
48	Tumor necrosis factor region polymorphisms are associated with AIDS and with cytomegalovirus retinitis. Aids, 2009, 23, 1641-1647.	1.0	16
49	The genetic diversity within the $1.4\hat{a}$ \in %kb HLA-G $5\hat{a}$ \in 2 upstream regulatory region moderately impacts on cellular microenvironment responses. Scientific Reports, 2018, 8, 5652.	1.6	16
50	SNPâ€HLA Reference Consortium (SHLARC): HLA and SNP data sharing for promoting MHC entric analyses in genomics. Genetic Epidemiology, 2020, 44, 733-740.	0.6	16
51	Human Leukocyte Antigen-G: A Promising Prognostic Marker of Disease Progression to Improve the Control of Human African Trypanosomiasis. Clinical Infectious Diseases, 2016, 63, ciw505.	2.9	15
52	Current HLA Investigations on SARS-CoV-2 and Perspectives. Frontiers in Genetics, 2021, 12, 774922.	1.1	15
53	HLA-E regulatory and coding region variability and haplotypes in a Brazilian population sample. Molecular Immunology, 2017, 91, 173-184.	1.0	14
54	HLA-G, -E and -F regulatory and coding region variability and haplotypes in the Beninese Toffin population sample. Molecular Immunology, 2018, 104, 108-127.	1.0	14

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55	How HLA diversity is apportioned: influence of selection and relevance to transplantation. Philosophical Transactions of the Royal Society B: Biological Sciences, 2022, 377, 20200420.	1.8	14
56	The +3187A/G HLA $\hat{a}\in G$ polymorphic site is associated with polar forms and reactive reaction in leprosy. Molecular Genetics & Enomic Medicine, 2013, 1, 123-130.	0.6	12
57	<scp><i>HLA </i></scp> genetic diversity and evolutionary insights in two samples from Brazil and Benin. Hla, 2020, 96, 468-486.	0.4	12
58	Insights on hair, skin and eye color of ancient and contemporary Native Americans. Forensic Science International: Genetics, 2020, 48, 102335.	1.6	12
59	Approaching Genetics Through the MHC Lens: Tools and Methods for HLA Research. Frontiers in Genetics, 2021, 12, 774916.	1.1	12
60	Interferon- \hat{l}^3 +874 Polymorphism in the First Intron of the Human Interferon- \hat{l}^3 Gene and Kidney Allograft Outcome. Transplantation Proceedings, 2010, 42, 4505-4508.	0.3	11
61	HLA-G genetic diversity and evolutive aspects in worldwide populations. Scientific Reports, 2021, 11, 23070.	1.6	11
62	HLAâ€G polymorphism and breast cancer. International Journal of Immunogenetics, 2014, 41, 143-148.	0.8	10
63	<i>HLAâ€A</i> promoter, coding, and 3′UTR sequences in a Brazilian cohort, and their evolutionary aspects. Hla, 2019, 93, 65-79.	0.4	10
64	An immunogenetic view of COVID-19. Genetics and Molecular Biology, 2021, 44, e20210036.	0.6	10
65	A novel HLAâ€G allele, <i>HLAâ€G*010111</i> , in the Brazilian population. Tissue Antigens, 2007, 70, 349-350.	1.0	9
66	Human leukocyte antigen-G 3' untranslated region polymorphisms are associated with asthma severity. Molecular Immunology, 2018, 101, 500-506.	1.0	9
67	Extended HLA-G genetic diversity and ancestry composition in a Brazilian admixed population sample: Implications for HLA-G transcriptional control and for case-control association studies. Human Immunology, 2018, 79, 790-799.	1.2	9
68	A large familial cluster and sporadic cases of frontal fibrosing alopecia in Brazil reinforce known human leucocyte antigen (HLA) associations and indicate new HLA susceptibility haplotypes. Journal of the European Academy of Dermatology and Venereology, 2020, 34, 2409-2413.	1.3	9
69	Absence of the HLA-G*0113N allele in Amerindian populations from the Brazilian Amazon region. Human Immunology, 2010, 71, 428-431.	1.2	8
70	P014 HLA-C, HLA-E and HLA-G regulatory and coding region polymorphisms in patients exhibiting gestational diabetes mellitus. Human Immunology, 2017, 78, 64.	1.2	8
71	Applicability of the SNPforID 52-plex panel for human identification and ancestry evaluation in a Brazilian population sample by next-generation sequencing. Forensic Science International: Genetics, 2019, 40, 201-209.	1.6	8
72	Prediction of eye and hair pigmentation phenotypes using the HIrisPlex system in a Brazilian admixed population sample. International Journal of Legal Medicine, 2021, 135, 1329-1339.	1.2	8

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73	Manifesting carriers of X-linked myotubular myopathy. Neurology: Genetics, 2020, 6, e513.	0.9	7
74	Identification of a novel 120 bp allele at the TNFd microsatellite locus. Tissue Antigens, 2006, 67, 318-320.	1.0	6
75	A nonsynonymous mutation at <i><scp>HLA</scp>â€E</i> defines the new <i>E*01:06</i> allele in Brazilian individuals. Tissue Antigens, 2013, 82, 216-217.	1.0	6
76	Analysis and comparison of the STR genotypes called with HipSTR, STRait Razor and toaSTR by using next generation sequencing data in a Brazilian population sample. Forensic Science International: Genetics, 2022, 58, 102676.	1.6	6
77	Genetic diversity of the <i>LILRB1</i> and <i>LILRB2</i> coding regions in an admixed Brazilian population sample. Hla, 2022, 100, 325-348.	0.4	6
78	Typing class I HLA-A gene using a nested PCR-RFLP procedure. Brazilian Journal of Medical and Biological Research, 2005, 38, 837-842.	0.7	5
79	Evaluation of MC1R high-throughput nucleotide sequencing data generated by the 1000 Genomes Project. Genetics and Molecular Biology, 2017, 40, 530-539.	0.6	5
80	HLA-G liver expression and HLA-G extended haplotypes are associated with chronic hepatitis C in HIV-negative and HIV-coinfected patients. Clinical Immunology, 2020, 217, 108482.	1.4	5
81	Recurrence of COVID-19 associated with reduced T-cell responses in a monozygotic twin pair. Open Biology, 2022, 12, 210240.	1.5	5
82	Identification of two new <i>HLAâ€G</i> alleles, <i>G*01:01:03:03</i> and <i>G*01:01:21</i> , in Brazilian individuals. Tissue Antigens, 2012, 80, 70-71.	1.0	4
83	Lack of association between <i><scp>HLA</scp>â€E</i> polymorphisms and transitional cell carcinoma of the bladder. Tissue Antigens, 2013, 82, 197-200.	1.0	4
84	Low variability at the HLA-E promoter region in the Brazilian population. Human Immunology, 2016, 77, 172-175.	1.2	4
85	HLA-C Alleles and Cytomegalovirus Retinitis in Brazilian Patients with AIDS. Journal of Ophthalmology, 2018, 2018, 1-5.	0.6	4
86	TNF microsatellite alleles may confer protection against the development of lipodystrophy syndrome in Brazilian HIV patients. International Journal of Immunogenetics, 2010, 37, 379-385.	0.8	3
87	Simple Methods for the Detection of HLA-G Variants in Coding and Non-coding Regions. Methods in Molecular Biology, 2012, 882, 123-142.	0.4	3
88	A novel HLAâ€E allele, <i>E*01:03:05</i> , identified in two Brazilian individuals. Tissue Antigens, 2012, 80, 200-201.	1.0	3
89	Relevance of PD-L1 Non-Coding Polymorphisms on the Prognosis of a Genetically Admixed NSCLC Cohort. Pharmacogenomics and Personalized Medicine, 2021, Volume 14, 239-252.	0.4	3
90	HLA-E gene polymorphisms in chronic hepatitis C: Impact on HLA-E liver expression and disease severity. Human Immunology, 2021, 82, 177-185.	1.2	3

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91	Evaluation of the HIrisPlex-S system in a Brazilian population sample. Forensic Science International: Genetics Supplement Series, 2019, 7, 794-796.	0.1	2
92	Variability at the 3′ untranslated region of the HLA-G gene: a study on patients with AIDS and cytomegalovirus retinochoroiditis. Scientific Reports, 2020, 10, 18646.	1.6	1
93	KIR2DL4 genetic diversity in a Brazilian population sample: implications for transcription regulation and protein diversity in samples with different ancestry backgrounds. Immunogenetics, 2021, 73, 227-241.	1.2	1
94	Peripheral spectrum neurological disorder after arbovirus infection is associated with HLA-F variants among Northeastern Brazilians. Infection, Genetics and Evolution, 2021, 92, 104855.	1.0	1
95	88-P: HLA-G 5' Regulatory Region and 3' Untranslated Region Variability in the Brazilian Population. Human Immunology, 2010, 71, S76.	1.2	O
96	599 ALLELES AND GENOTYPES OF IL-18 AND TNF-α PROMOTER POLYMORPHISMS ARE ASSOCIATED WITH HIGHER RISKOF HEPATOCELLULAR CARCINOMA (HCC) IN BRAZILIAN POPULATION. Journal of Hepatology, 2010, 52, S236.	1.8	0
97	713 ASSOCIATION OF INTERLEUKIN-18 (IL-18), TUMOR NECROSIS FACTOR-α (TNF-α) AND INTERFERON-γ (IFN-γ GENES POLYMORPHISMS WITH SEVERITY OF LIVER FIBROSIS IN BRAZILIAN PATIENTS WITH CHRONIC HEPATITIS B. Journal of Hepatology, 2010, 52, S277.	1.8	O
98	Genetic Analysis As a Practical Tool to Diagnose Hereditary Angioedema with Normal C1 Inhibitor: A Case Report. Journal of Allergy and Clinical Immunology, 2015, 135, AB197.	1.5	0
99	Human leukocyte antigen (HLA)-F and -G gene polymorphisms and haplotypes are associated with malaria susceptibility in the Beninese Toffin children. Infection, Genetics and Evolution, 2021, 92, 104828.	1.0	O
100	Genetic Analysis as a Practical Tool for Diagnosis of Hereditary Angioedema With Normal C1 Inhibitor: A Case Report. Journal of Investigational Allergology and Clinical Immunology, 2016, 26, 48-72.	0.6	0