Rafael De Cid

List of Publications by Year in descending order

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80 papers 5,876 citations

30 h-index 79698 73 g-index

89 all docs 89 docs citations

89 times ranked 12539 citing authors

#	Article	IF	CITATIONS
1	Genomewide Association Study of Severe Covid-19 with Respiratory Failure. New England Journal of Medicine, 2020, 383, 1522-1534.	27.0	1,548
2	The COVID-19 Host Genetics Initiative, a global initiative to elucidate the role of host genetic factors in susceptibility and severity of the SARS-CoV-2 virus pandemic. European Journal of Human Genetics, 2020, 28, 715-718.	2.8	649
3	Deletion of the late cornified envelope LCE3B and LCE3C genes as a susceptibility factor for psoriasis. Nature Genetics, 2009, 41, 211-215.	21.4	482
4	Brain-Derived Neurotrophic Factor Val66Met and Psychiatric Disorders: Meta-Analysis of Case-Control Studies Confirm Association to Substance-Related Disorders, Eating Disorders, and Schizophrenia. Biological Psychiatry, 2007, 61, 911-922.	1.3	381
5	Identification of SLC7A7, encoding y+LAT-1, as the lysinuric protein intolerance gene. Nature Genetics, 1999, 21, 293-296.	21.4	286
6	Met66 in the brain-derived neurotrophic factor (BDNF) precursor is associated with anorexia nervosa restrictive type. Molecular Psychiatry, 2003, 8, 745-751.	7.9	176
7	Identification of MYH Mutation Carriers in Colorectal Cancer: A Multicenter, Case-Control, Population-Based Study. Clinical Gastroenterology and Hepatology, 2007, 5, 379-387.	4.4	141
8	Coding haplotype analysis supports HCR as the putative susceptibility gene for psoriasis at the MHC PSORS1 locus. Human Molecular Genetics, 2002, 11, 589-597.	2.9	131
9	Identification of new putative susceptibility genes for several psychiatric disorders by association analysis of regulatory and nonâ€synonymous SNPs of 306 genes involved in neurotransmission and neurodevelopment. American Journal of Medical Genetics Part B: Neuropsychiatric Genetics, 2009, 150B. 808-816.	1.7	98
10	DNA Hypomethylation at ALOX12Is Associated with Persistent Wheezing in Childhood. American Journal of Respiratory and Critical Care Medicine, 2012, 185, 937-943.	5.6	97
11	Meta-Analysis Confirms the LCE3C_LCE3B Deletion as a Risk Factor for Psoriasis in Several Ethnic Groups and Finds Interaction with HLA-Cw6. Journal of Investigative Dermatology, 2011, 131, 1105-1109.	0.7	89
12	Association of Early-life Exposure to Household Gas Appliances and Indoor Nitrogen Dioxide With Cognition and Attention Behavior in Preschoolers. American Journal of Epidemiology, 2009, 169, 1327-1336.	3.4	81
13	Traffic-Related Air Pollution, Oxidative Stress Genes, and Asthma (ECHRS). Environmental Health Perspectives, 2009, 117, 1919-1924.	6.0	78
14	A brain-derived neurotrophic factor (BDNF) haplotype is associated with antidepressant treatment outcome in mood disorders. Pharmacogenomics Journal, 2008, 8, 101-112.	2.0	76
15	Altered brainâ€derived neurotrophic factor blood levels and gene variability are associated with anorexia and bulimia. Genes, Brain and Behavior, 2007, 6, 706-716.	2.2	7 3
16	Extensive Genotyping of the BDNF and NTRK2 Genes Define Protective Haplotypes Against Obsessive-Compulsive Disorder. Biological Psychiatry, 2008, 63, 619-628.	1.3	66
17	SNP analysis to results (SNPator): a web-based environment oriented to statistical genomics analyses upon SNP data. Bioinformatics, 2008, 24, 1643-1644.	4.1	61
18	Ambient Air Pollution in Relation to SARS-CoV-2 Infection, Antibody Response, and COVID-19 Disease: A Cohort Study in Catalonia, Spain (COVICAT Study). Environmental Health Perspectives, 2021, 129, 117003.	6.0	58

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19	<i>BDNF </i> variability in opioid addicts and response to methadone treatment: preliminary findings. Genes, Brain and Behavior, 2008, 7, 515-522.	2.2	53
20	Missense mutations in the cystic fibrosis gene in adult patients with asthma. Human Mutation, 1999, 14, 510-519.	2.5	51
21	Association of NTRK3 and its interaction with NGF suggest an altered cross-regulation of the neurotrophin signaling pathway in eating disorders. Human Molecular Genetics, 2008, 17, 1234-1244.	2.9	50
22	Detection of unrecognized low-level mtDNA heteroplasmy may explain the variable phenotypic expressivity of apparently homoplasmic mtDNA mutations. Human Mutation, 2008, 29, 248-257.	2.5	44
23	A new titinopathy. Neurology, 2015, 85, 2126-2135.	1.1	44
24	Genetic susceptibility to obsessiveâ€compulsive hoarding: the contribution of neurotrophic tyrosine kinase receptor type 3 gene ¹ . Genes, Brain and Behavior, 2008, 7, 778-785.	2.2	43
25	Joint effect of obesity and TNFA variability on asthma: two international cohort studies. European Respiratory Journal, 2009, 33, 1003-1009.	6.7	43
26	A pooling-based genome-wide analysis identifies new potential candidate genes for atopy in the European Community Respiratory Health Survey (ECRHS). BMC Medical Genetics, 2009, 10, 128.	2.1	43
27	Exploring the Role of Mutations in Fanconi Anemia Genes in Hereditary Cancer Patients. Cancers, 2020, 12, 829.	3.7	41
28	Association Study of the Chromosomal Region Containing the FCER2 Gene Suggests It Has a Regulatory Role in Atopic Disorders. American Journal of Respiratory and Critical Care Medicine, 2000, 161, 700-706.	5.6	40
29	A new approach for identifying non-pathogenic mutations. An analysis of the cystic fibrosis transmembrane regulator gene in normal individuals. Human Genetics, 2000, 106, 172-178.	3.8	39
30	Splice-site mutation in the PDS gene may result in intrafamilial variability for deafness in Pendred syndrome., 1999, 14, 520-526.		37
31	A Brain-Derived Neurotrophic Factor Haplotype Is Associated with Therapeutic Response in Obsessive-Compulsive Disorder. Biological Psychiatry, 2009, 66, 674-680.	1.3	34
32	Influence of Glutathione S -Transferase Polymorphisms on Cognitive Functioning Effects Induced by p,p ′-DDT among Preschoolers. Environmental Health Perspectives, 2008, 116, 1581-1585.	6.0	32
33	Deletion of LCE3C and LCE3B is a susceptibility factor for psoriatic arthritis: A study in Spanish and Italian populations and meta-analysis. Arthritis and Rheumatism, 2011, 63, 1860-1865.	6.7	31
34	GCAT Genomes for life: a prospective cohort study of the genomes of Catalonia. BMJ Open, 2018, 8, e018324.	1.9	31
35	Independent Contribution of Common CFTR Variants to Chronic Pancreatitis. Pancreas, 2010, 39, 209-215.	1.1	30
36	TNFA -308G>A in two international population-based cohorts and risk of asthma. European Respiratory Journal, 2008, 32, 350-361.	6.7	28

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37	An Association Study of 22 Candidate Genes in Psoriasis Families Reveals Shared Genetic Factors with Other Autoimmune and Skin Disorders. Journal of Investigative Dermatology, 2009, 129, 2637-2645.	0.7	28
38	Response to Methadone Maintenance Treatment is Associated with the MYOCD and GRM6 Genes. Molecular Diagnosis and Therapy, 2010, 14, 171-178.	3.8	28
39	Multitrait genome association analysis identifies new susceptibility genes for human anthropometric variation in the GCAT cohort. Journal of Medical Genetics, 2018, 55, 765-778.	3.2	28
40	ADAR1 affects HCV infection by modulating innate immune response. Antiviral Research, 2018, 156, 116-127.	4.1	27
41	Identification of five new mutations of PDS/SLC26A4 in Mediterranean families with hearing impairment. Human Mutation, 2001, 18, 548-548.	2.5	26
42	GSTM1 polymorphisms modify the effect of maternal smoking during pregnancy on cognitive functioning in preschoolers. International Journal of Epidemiology, 2009, 38, 690-697.	1.9	26
43	Immunological Self-Tolerance: An Analysis Employing Cytokines or Cytokine Receptors Encoded by Transgenes or a Recombinant Vaccinia Virus. Immunological Reviews, 1991, 122, 173-204.	6.0	25
44	Positionally cloned genes and age-specific effects in asthma and atopy: an international population-based cohort study (ECRHS). Thorax, 2010, 65, 124-131.	5.6	25
45	An Autosomal-Recessive Form of Cutis Laxa Is Due to Homozygous Elastin Mutations, and the Phenotype May Be Modified by a Heterozygous Fibulin 5 Polymorphism. Journal of Investigative Dermatology, 2009, 129, 1650-1655.	0.7	24
46	CFTR and asthma in the French EGEA study. European Journal of Human Genetics, 2001, 9, 67-69.	2.8	23
47	Gene-environment interactions in asthma. Occupational and Environmental Medicine, 2006, 63, 776-786.	2.8	22
48	Disease networks identify specific conditions and pleiotropy influencing multimorbidity in the general population. Scientific Reports, 2018, 8, 15970.	3.3	22
49	Detection of TRIM32 deletions in LGMD patients analyzed by a combined strategy of CGH array and massively parallel sequencing. European Journal of Human Genetics, 2015, 23, 929-934.	2.8	21
50	Functional double-negative T cells in the periphery express T cell receptor \hat{V}^2 gene products that cause deletion of single-positive T cells. European Journal of Immunology, 1993, 23, 250-254.	2.9	19
51	Absence of cytogenetic effects in children and adults with attention-deficit/hyperactivity disorder treated with methylphenidate. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2009, 666, 44-49.	1.0	18
52	Association of the ARLTS1 Cys148Arg variant with sporadic and familial colorectal cancer. Carcinogenesis, 2007, 28, 1687-1691.	2.8	16
53	Infection induced SARS-CoV-2 seroprevalence and heterogeneity of antibody responses in a general population cohort study in Catalonia Spain. Scientific Reports, 2021, 11, 21571.	3.3	16
54	Assessment of kinship detection using RNA-seq data. Nucleic Acids Research, 2019, 47, e136-e136.	14.5	13

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55	Founder effect of a pathogenic <i>MSH2</i> mutation identified in Spanish families with Lynch syndrome. Clinical Genetics, 2010, 78, 186-190.	2.0	11
56	A Log-Ratio Biplot Approach for Exploring Genetic Relatedness Based on Identity by State. Frontiers in Genetics, 2019, 10, 341.	2.3	11
57	BARD1 Pathogenic Variants Are Associated with Triple-Negative Breast Cancer in a Spanish Hereditary Breast and Ovarian Cancer Cohort. Genes, 2021, 12, 150.	2.4	11
58	Molecular characterization of a $t(9;12)(p21;q13)$ balanced chromosome translocation in combination with integrative genomics analysis identifies C9 or f14 as a candidate tumor-suppressor. Genes Chromosomes and Cancer, 2007, 46, 155-162.	2.8	10
59	Role of the neurotrophin network in eating disorders' subphenotypes: Body mass index and age at onset of the disease. Journal of Psychiatric Research, 2010, 44, 834-840.	3.1	10
60	ALDH5A1 variability in opioid dependent patients could influence response to methadone treatment. European Neuropsychopharmacology, 2014, 24, 420-424.	0.7	10
61	Comprehensive analysis and ACMGâ€based classification of <i>CHEK2</i> variants in hereditary cancer patients. Human Mutation, 2020, 41, 2128-2142.	2.5	10
62	Genomic profiling in advanced stage non-small-cell lung cancer patients with platinum-based chemotherapy identifies germline variants with prognostic value in SMYD2. Cancer Treatment and Research Communications, 2018, 15, 21-31.	1.7	9
63	Association study of proposed candidate genes/regions in a population of Spanish asthmatics. European Journal of Epidemiology, 2000, 16, 745-750.	5.7	8
64	Early life environment, neurodevelopment and the interrelation with atopy. Environmental Research, 2010, 110, 733-738.	7. 5	8
65	ADAR1 function affects HPV replication and is associated to recurrent human papillomavirus-induced dysplasia in HIV coinfected individuals. Scientific Reports, 2019, 9, 19848.	3.3	8
66	ERCC3, a new ovarian cancer susceptibility gene?. European Journal of Cancer, 2020, 141, 1-8.	2.8	8
67	Revisiting the UK Genetic Severity Score for NF2: a proposal for the addition of a functional genetic component. Journal of Medical Genetics, 2022, 59, 678-686.	3.2	7
68	Immune Cell Associations with Cancer Risk. IScience, 2020, 23, 101296.	4.1	6
69	Single nucleotide polymorphisms in PNPLA3, ADAR-1 and IFIH1 are associated with advanced liver fibrosis in patients co-infected with HIV-1//hepatitis C virus. Aids, 2021, 35, 2497-2502.	2.2	6
70	GCAT Panel, a comprehensive structural variant haplotype map of the Iberian population from high-coverage whole-genome sequencing. Nucleic Acids Research, 2022, 50, 2464-2479.	14.5	6
71	A likelihood ratio approach for identifying three-quarter siblings in genetic databases. Heredity, 2021, 126, 537-547.	2.6	5
72	Association of NTRK3 and its interaction with NGF suggest an altered cross-regulation of the neurotrophin signaling pathway in eating disorders. European Psychiatry, 2008, 23, S182.	0.2	2

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73	Pipeline design to identify key features and classify the chemotherapy response on lung cancer patients using large-scale genetic data. BMC Systems Biology, 2018, 12, 97.	3.0	2
74	Ambient air pollution and risk of SARS-CoV-2 infection and of COVID-19 disease in a cohort study in Catalonia (COVICAT Cohort). ISEE Conference Abstracts, 2021, 2021, .	0.0	2
75	G.P.281. Neuromuscular Disorders, 2014, 24, 901-902.	0.6	O
76	Exome sequencing identifies novel truncating TTN mutations with Emery–Dreifuss like muscular dystrophy and secondary calpain3 deficiency without cardiac abnormality. Neuromuscular Disorders, 2015, 25, S245.	0.6	0
77	SNP Selection for Psychiatric Disease Association Based on Allele Frequency Plots. , 2009, , .		0
78	Evidence for shared genetic risk factors between lymphangioleiomyomatosis and pulmonary function. ERJ Open Research, 2022, 8, 00375-2021.	2.6	0
79	O-446â€Occupational risk factors for SARS-CoV-2 infection and COVID-19: Results from the COVICAT cohort study in Catalonia, Spain , 2021, , .		0
80	RNA assay identifies a previous misclassification of BARD1 c.1977A>G variant. Scientific Reports, 2021, 11, 22948.	3.3	0