## Luis M. Real

## List of Publications by Year in descending order

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116 papers	6,134 citations	33 h-index	95266 68 g-index
123	123	123	9553
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	Genetic meta-analysis of diagnosed Alzheimer's disease identifies new risk loci and implicates Aβ, tau, immunity and lipid processing. Nature Genetics, 2019, 51, 414-430.	21.4	1,962
2	New insights into the genetic etiology of Alzheimer's disease and related dementias. Nature Genetics, 2022, 54, 412-436.	21.4	700
3	Galectin-3, a novel endogenous TREM2 ligand, detrimentally regulates inflammatory response in Alzheimer's disease. Acta Neuropathologica, 2019, 138, 251-273.	7.7	187
4	Common variants in Alzheimer's disease and risk stratification by polygenic risk scores. Nature Communications, 2021, 12, 3417.	12.8	140
5	CCR5 Expression Influences the Progression of Human Breast Cancer in a p53-dependent Manner. Journal of Experimental Medicine, 2003, 198, 1381-1389.	8.5	129
6	Role of follicle-stimulating hormone receptor Ser680Asn polymorphism in the efficacy of follicle-stimulating hormone. Fertility and Sterility, 2003, 80, 571-576.	1.0	122
7	Analysis of HLA class I expression in progressing and regressing metastatic melanoma lesions after immunotherapy. Immunogenetics, 2008, 60, 439-447.	2.4	119
8	Genomeâ€wide association analysis of dementia and its clinical endophenotypes reveal novel loci associated with Alzheimer's disease and three causality networks: The GR@ACE project. Alzheimer's and Dementia, 2019, 15, 1333-1347.	0.8	111
9	Human controlled ovarian hyperstimulation outcome is a polygenic trait. Pharmacogenetics and Genomics, 2004, 14, 285-293.	5.7	109
10	Chromosome loss is the most frequent mechanism contributing to HLA haplotype loss in human tumors., 1999, 83, 91-97.		104
11	The membrane-spanning 4-domains, subfamily A (MS4A) gene cluster contains a common variant associated with Alzheimer's disease. Genome Medicine, 2011, 3, 33.	8.2	81
12	HLA class I expression in metastatic melanoma correlates with tumor development during autologous vaccination. Cancer Immunology, Immunotherapy, 2007, 56, 709-717.	4.2	78
13	CAPN10 Alleles Are Associated with Polycystic Ovary Syndrome. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 3971-3976.	3.6	76
14	A method for detecting epistasis in genome-wide studies using case-control multi-locus association analysis. BMC Genomics, 2008, 9, 360.	2.8	76
15	Expression of HLA G in human tumors is not a frequent event. , 1999, 81, 512-518.		65
16	Specific CAPN10 Gene Haplotypes Influence the Clinical Profile of Polycystic Ovary Patients. Journal of Clinical Endocrinology and Metabolism, 2003, 88, 5529-5536.	3.6	63
17	Unresponsiveness to interferon associated with STAT1 protein deficiency in a gastric adenocarcinoma cell line. Cancer Immunology, Immunotherapy, 1998, 47, 113-120.	4.2	62
18	ATP5H/KCTD2 locus is associated with Alzheimer's disease risk. Molecular Psychiatry, 2014, 19, 682-687.	7.9	62

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19	Bone morphogenetic protein 15 (BMP15) alleles predict over-response to recombinant follicle stimulation hormone and iatrogenic ovarian hyperstimulation syndrome (OHSS). Pharmacogenetics and Genomics, 2006, 16, 485-495.	1.5	58
20	Regressing and progressing metastatic lesions: resistance to immunotherapy is predetermined by irreversible HLA class I antigen alterations. Cancer Immunology, Immunotherapy, 2008, 57, 1727-1733.	4.2	56
21	Incidence and natural history of hepatitis E virus coinfection among HIV-infected patients. Aids, 2014, 28, 1931-1937.	2.2	52
22	Increased risk of severe clinical course of COVID-19 in carriers of HLA-C*04:01. EClinicalMedicine, 2021, 40, 101099.	7.1	52
23	Association of NOS3 gene with metabolic syndrome in hypertensive patients. Thrombosis and Haemostasis, 2004, 92, 413-418.	3.4	50
24	Activation of PKR Causes Amyloid ß-Peptide Accumulation via De-Repression of BACE1 Expression. PLoS ONE, 2011, 6, e21456.	2.5	50
25	Genetic Structure of the Spanish Population. BMC Genomics, 2010, 11, 326.	2.8	49
26	Multilocus analyses of estrogen-related genes reveal involvement of the ESR1 gene in male infertility and the polygenic nature of the pathology. Fertility and Sterility, 2005, 84, 910-918.	1.0	47
27	Genome-Wide Association Study of Multiple Sclerosis Confirms a Novel Locus at 5p13.1. PLoS ONE, 2012, 7, e36140.	2.5	46
28	Multiple mechanisms of immune evasion can coexist in melanoma tumor cell lines derived from the same patient. Cancer Immunology, Immunotherapy, 2001, 49, 621-628.	4.2	45
29	Identification of genetic factors associated with susceptibility to angiotensin-converting enzyme inhibitors-induced cough. Pharmacogenetics and Genomics, 2011, 21, 10-17.	1.5	45
30	Hepatitis C virus reinfection after sustained virological response in HIV-infected patients with chronic hepatitis C. Journal of Infection, 2015, 71, 571-577.	3.3	42
31	Changes in Liver Steatosis After Switching From Efavirenz to Raltegravir Among Human Immunodeficiency Virus–Infected Patients With Nonalcoholic Fatty Liver Disease. Clinical Infectious Diseases, 2017, 65, 1012-1019.	5.8	42
32	Effects of first-line antiretroviral therapy on the CD4/CD8 ratio and CD8 cell counts in CoRIS: a prospective multicentre cohort study. Lancet HIV, the, 2020, 7, e565-e573.	4.7	42
33	Association of genetic markers within the KIT and KITLG genes with human male infertility. Human Reproduction, 2006, 21, 3185-3192.	0.9	40
34	CALHM1 P86L Polymorphism is Associated with Late-Onset Alzheimer's Disease in a Recessive Model. Journal of Alzheimer's Disease, 2010, 20, 247-251.	2.6	38
35	Genetic Study of Neurexin and Neuroligin Genes in Alzheimer's Disease. Journal of Alzheimer's Disease, 2013, 35, 403-412.	2.6	38
36	Differential effect on U937 cell differentiation by targeting transcriptional factors implicated in tissue- or stage-specific induced integrin expression. Experimental Hematology, 1999, 27, 353-364.	0.4	37

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37	Estrogen receptor alpha gene variants are associated with Alzheimer's disease. Neurobiology of Aging, 2012, 33, 198.e15-198.e24.	3.1	36
38	In vivo and in vitro generation of a new altered HLA phenotype in melanoma-tumour-cell variants expressing a single HLA-class-I allele., 1998, 75, 317-323.		31
39	Association of Rare <i>APOE</i> Missense Variants V236E and R251G With Risk of Alzheimer Disease. JAMA Neurology, 2022, 79, 652.	9.0	31
40	CAPN10 Alleles Are Associated with Polycystic Ovary Syndrome. Journal of Clinical Endocrinology and Metabolism, 2002, 87, 3971-3976.	3.6	30
41	A Colorectal Cancer Susceptibility New Variant at 4q26 in the Spanish Population Identified by Genome-Wide Association Analysis. PLoS ONE, 2014, 9, e101178.	2.5	26
42	Pharmacogenetics of controlled ovarian hyperstimulation. Pharmacogenomics, 2005, 6, 629-637.	1.3	25
43	WWOX gene is associated with HDL cholesterol and triglyceride levels. BMC Medical Genetics, 2010, 11, 148.	2.1	24
44	IFNL4 ss469415590 Variant Shows Similar Performance to rs12979860 as Predictor of Response to Treatment against Hepatitis C Virus Genotype 1 or 4 in Caucasians. PLoS ONE, 2014, 9, e95515.	2.5	24
45	Reassessment of Genotype 1 Hepatitis C Virus Subtype Misclassification by LiPA 2.0: Implications for Direct-Acting Antiviral Treatment. Journal of Clinical Microbiology, 2014, 52, 4027-4029.	3.9	23
46	Changes in liver steatosis evaluated by transient elastography with the controlled attenuation parameter in <scp>HIV</scp> â€infected patients. HIV Medicine, 2016, 17, 766-773.	2.2	23
47	Microsatellite instability analysis in tumors with different mechanisms for total loss of HLA expression. Cancer Immunology, Immunotherapy, 2000, 48, 684-690.	4.2	21
48	Lack of Association Between NOS3 Glu298Asp and Breast Cancer Risk: a Case–ontrol Study. Breast Cancer Research and Treatment, 2006, 100, 331-333.	2.5	21
49	Specific haplotypes of the CALPAIN-5 gene are associated with polycystic ovary syndrome. Human Reproduction, 2006, 21, 943-951.	0.9	20
50	Scanning of Y-chromosome azoospermia factors loci using real-time polymerase chain reaction and melting curve analysis. Fertility and Sterility, 2003, 80, 907-913.	1.0	19
51	Impact of genetic polymorphisms associated with nonalcoholic fatty liver disease on HIV-infected individuals. Aids, 2015, 29, 1927-1935.	2.2	19
52	IFNL4 rs368234815 polymorphism is associated with innate resistance to HIV-1 infection. Aids, 2015, 29, 1895-1897.	2.2	19
53	Genetic analysis of caveolin-1 and eNOS genes in colorectal cancer. Oncology Reports, 2006, 16, 353-9.	2.6	18
54	Genetic analysis of the GRM1 gene in human melanoma susceptibility. European Journal of Human Genetics, 2007, 15, 1176-1182.	2.8	17

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55	The Ubiquitin Proteasome System in Neuromuscular Disorders: Moving Beyond Movement. International Journal of Molecular Sciences, 2020, 21, 6429.	4.1	17
56	<scp>CCR</scp> 5 deficiency impairs <scp>CD</scp> 4 <sup>+</sup> Tâ€eell memory responses and antigenic sensitivity through increased ceramide synthesis. EMBO Journal, 2020, 39, e104749.	7.8	17
57	Genetic association of complement receptor 1 polymorphism rs3818361 in Alzheimer's disease. Alzheimer's and Dementia, 2011, 7, e124-9.	0.8	16
58	Association of low-density lipoprotein receptor genotypes with hepatitis C viral load. Genes and Immunity, 2014, 15, 16-24.	4.1	16
59	Sustained virological response to direct-acting antiviral regimens reduces the risk of hepatocellular carcinoma in HIV/HCV-coinfected patients with cirrhosis. Journal of Antimicrobial Chemotherapy, 2018, 73, 2435-2443.	3.0	16
60	Preliminary molecular genetic analysis of the Receptor Interacting Protein 140 (RIP140) in women affected by endometriosis. Journal of Experimental & Clinical Assisted Reproduction, 2005, 2, 11.	0.4	15
61	A Digenic Combination of Polymorphisms Within ESR1 and ESR2 Genes Are Associated With Age at Menarche in the Spanish Population. Reproductive Sciences, 2008, 15, 305-311.	2.5	15
62	Exploratory analysis of seven Alzheimer's disease genes: disease progression. Neurobiology of Aging, 2013, 34, 1310.e1-1310.e7.	3.1	15
63	The PNPLA3 Genetic Variant rs738409 Influences the Progression to Cirrhosis in HIV/Hepatitis C Virus Coinfected Patients. PLoS ONE, 2016, 11, e0168265.	2.5	15
64	HIV infection does not increase the risk of liver complications in hepatitis C virus-infected patient with advanced fibrosis, after sustained virological response with direct-acting antivirals. Aids, 2019, 33, 1167-1174.	2.2	15
65	Multiomics integrative analysis identifies APOE allele-specific blood biomarkers associated to Alzheimer's disease etiopathogenesis. Aging, 2021, 13, 9277-9329.	3.1	15
66	Identification of a protective haplogenotype within CAPN10 gene influencing colorectal cancer susceptibility. Journal of Gastroenterology and Hepatology (Australia), 2007, 22, 2298-2302.	2.8	14
67	HERC1 Ubiquitin Ligase Is Required for Normal Axonal Myelination in the Peripheral Nervous System. Molecular Neurobiology, 2018, 55, 8856-8868.	4.0	14
68	Looking for HLA-G expression in human tumours. Journal of Reproductive Immunology, 1999, 43, 263-273.	1.9	13
69	Re: Polymorphisms Associated With Circulating Sex Hormone Levels in Postmenopausal Women. Journal of the National Cancer Institute, 2005, 97, 152-153.	6.3	13
70	Whole-genome conditional two-locus analysis identifies novel candidate genes for late-onset Parkinson's disease. Neurogenetics, 2009, 10, 173-181.	1.4	13
71	<i>Calpain 10</i> gene and laryngeal cancer: A survival analysis. Head and Neck, 2011, 33, 72-76.	2.0	13
72	A Regulatory Polymorphism in HAVCR2 Modulates Susceptibility to HIV-1 Infection. PLoS ONE, 2014, 9, e106442.	2.5	13

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73	Association of complement receptor 2 polymorphisms with innate resistance to HIV-1 infection. Genes and Immunity, 2015, 16, 134-141.	4.1	13
74	Analysis of the ERalpha germline Pvull marker in breast cancer risk. Medical Science Monitor, 2008, 14, CR136-43.	1.1	13
75	Non-Alcoholic Fatty Liver Disease in HIV Infection. AIDS Reviews, 2017, 19, 35-46.	1.0	12
76	High frequency of potential interactions between directâ€acting antivirals and concomitant therapy in <scp>HIV</scp> /hepatitis C virusâ€coinfected patients in clinical practice. HIV Medicine, 2017, 18, 445-451.	2.2	11
77	MicroRNA Profile of HCV Spontaneous Clarified Individuals, Denotes Previous HCV Infection. Journal of Clinical Medicine, 2019, 8, 849.	2.4	11
78	Detection of Pvull Polymorphism within Intron 1 of ESR1 Gene by Real-Time PCR. Clinical Chemistry and Laboratory Medicine, 2003, 41, 392-3.	2.3	10
79	CAPN10 alleles modify laryngeal cancer risk in the Spanish population. European Journal of Surgical Oncology, 2008, 34, 94-99.	1.0	10
80	Genetic analysis of candidate SNPs for metabolic syndrome in obstructive sleep apnea (OSA). Gene, 2013, 521, 150-154.	2.2	10
81	Human Immunodeficiency Virus (HIV) Infection Is Associated With Lower Risk of Hepatocellular Carcinoma After Sustained Virological Response to Direct-acting Antivirals in Hepatitis C Infected Patients With Advanced Fibrosis. Clinical Infectious Diseases, 2021, 73, e2109-e2116.	5.8	10
82	A polymorphism linked to <i><scp>RRAS</scp></i> , <i><scp>SCAF</scp>1</i> , <i><scp>IRF</scp>3</i> and <i><scp>BCL</scp>2L12</i> genes is associated with cirrhosis in hepatitis C virus carriers. Liver International, 2014, 34, 558-566.	3.9	9
83	Association of complement C3d receptor 2 genotypes with the acquisition of HIV infection in a trial of recombinant glycoprotein 120 vaccine. Aids, 2020, 34, 25-32.	2.2	9
84	Genetic analysis of CAV1 gene in hypertension and metabolic syndrome. Thrombosis and Haemostasis, 2006, 95, 696-701.	3.4	9
85	Absence of de novo Y-chromosome microdeletions in male children conceived through intracytoplasmic sperm injection. Fertility and Sterility, 2004, 82, 1679-1680.	1.0	8
86	Exploring allelic imbalance within paraffin-embedded tumor biopsies using pyrosequencing technology. Clinical Chemistry and Laboratory Medicine, 2006, 44, 1076-81.	2.3	7
87	Identification of a 2244 base pair interstitial deletion within the human ESR1 gene in the Spanish population. Journal of Medical Genetics, 2008, 45, 420-424.	3.2	7
88	GOLPH2 Gene Markers are Not Associated with Alzheimer's Disease in a Sample of the Spanish Population. Journal of Alzheimer's Disease, 2009, 18, 751-754.	2.6	7
89	Variations at multiple genes improve interleukin 28B genotype predictive capacity for response to therapy against hepatitis C infection. Aids, 2013, 27, 2715-2724.	2.2	7
90	Common haplotypes in CD209 promoter and susceptibility to HIV-1 infection in intravenous drug users. Infection, Genetics and Evolution, 2016, 45, 20-25.	2.3	7

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91	Similar incidence of coronavirus disease 2019 (COVID-19) in patients with rheumatic diseases with and without hydroxychloroquine therapy. PLoS ONE, 2021, 16, e0249036.	2.5	7
92	Genomic Characterization of Host Factors Related to SARS-CoV-2 Infection in People with Dementia and Control Populations: The GR@ACE/DEGESCO Study. Journal of Personalized Medicine, 2021, 11, 1318.	2.5	7
93	Fat mass and obesityâ€associated gene variations are related to fatty liver disease in <scp>HIV</scp> â€infected patients. HIV Medicine, 2017, 18, 546-554.	2.2	6
94	Long runs of homozygosity are associated with Alzheimer's disease. Translational Psychiatry, 2021, 11, 142.	4.8	6
95	Detection of Nucleotide c985 A→G Mutation of Medium-Chain Acyl-CoA Dehydrogenase Gene by Real-Time PCR. Clinical Chemistry, 2001, 47, 958-959.	3.2	5
96	Pyrosequencing protocol requiring a unique biotinylated primer. Clinical Chemistry and Laboratory Medicine, 2006, 44, 435-41.	2.3	5
97	The MTHFD1L Gene rs11754661 Marker is Not Associated with Alzheimer's Disease in a Sample of the Spanish Population. Journal of Alzheimer's Disease, 2011, 25, 47-50.	2.6	5
98	Baseline resistance-guided therapy does not enhance the response to interferon-free treatment of HCV infection in real life. Scientific Reports, 2018, 8, 14905.	3.3	4
99	A genomeâ€wide association study on low susceptibility to hepatitis C virus infection (GEHEP012 study). Liver International, 2019, 39, 1918-1926.	3.9	4
100	Liver Stiffness–Based Strategies Predict Absence of Variceal Bleeding in Cirrhotic Hepatitis C Virus–Infected Patients With and Without Human Immunodeficiency Virus Coinfection After Sustained Virological Response. Clinical Infectious Diseases, 2021, 72, e96-e102.	5.8	4
101	Kinetics of emergence of liver complications in hepatitis C virus infected patients and advanced fibrosis, with and without HIV-coinfection, after sustained virological response. Aids, 2021, 35, 2119-2127.	2.2	4
102	Lower probability of persistence of total anti-SARS-CoV-2 antibodies after COVID-19 among people living with HIV. Clinical Microbiology and Infection, 2022, , .	6.0	4
103	High efficacy of glecaprevir/pibrentasvir for HCV-infected individuals with active drug use. Journal of Infection, 2022, 85, 322-326.	3.3	4
104	Core amino acid variation at position 110 is associated with sustained virological response in Caucasian patients with chronic hepatitis C virus 1b infection. Archives of Virology, 2014, 159, 3345-3351.	2.1	3
105	Chromosome loss is the most frequent mechanism contributing to HLA haplotype loss in human tumors. International Journal of Cancer, 1999, 83, 91-97.	5.1	3
106	CD46 Genetic Variability and HIV-1 Infection Susceptibility. Cells, 2021, 10, 3094.	4.1	3
107	Molecular evaluation of human Ubiquilin 2 gene PXX domain in familial frontotemporal dementia patients. Journal of Neurology, 2012, 259, 2488-2490.	<b>3.</b> 6	2
108	Genetic markers of lipid metabolism genes associated with low susceptibility to HCV infection. Scientific Reports, 2019, 9, 9054.	3.3	2

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109	A Genome-Wide Association Study on Liver Stiffness Changes during Hepatitis C Virus Infection Cure. Diagnostics, 2021, 11, 1501.	2.6	2
110	Changes in the response to treatment against chronic hepatitis C between 1999 and 2015. European Journal of Gastroenterology and Hepatology, 2016, 28, 1253-1257.	1.6	1
111	Low incidence of acute hepatitis C virus infection among Southern Spanish HIV-infected individuals. Journal of Infection, 2017, 74, 514-517.	3.3	1
112	In vivo and in vitro generation of a new altered HLA phenotype in melanomaâ€tumourâ€cell variants expressing a single HLAâ€classâ€l allele. International Journal of Cancer, 1998, 75, 317-323.	5.1	1
113	Genetic Association Studies in Host–Pathogen Interaction Analysis. Methods in Molecular Biology, 2018, 1734, 1-11.	0.9	O
114	Impact of recent drug use on the efficacy of elbasvir/grazoprevir for HCVâ€infected people on opioid agonist therapy. Journal of Viral Hepatitis, 2021, 28, 878-886.	2.0	0
115	A polygenic risk score for mosaic loss of chromosome Y susceptibility is associated with higher risk of MCI to AD conversion Alzheimer's and Dementia, 2021, 17 Suppl 3, e053745.	0.8	O
116	<i>IFNL4</i> genotype influences the rate of HIV-1 seroconversion in men who have sex with men. Virulence, 2022, 13, 757-763.	4.4	O