Cinzia Ciccacci

List of Publications by Year
in descending order

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68	1,670	279487	315357
papers	citations	23 h-index	38 g-index

69	69	69	2685
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	CYP4F2genetic variant (rs2108622) significantly contributes to warfarin dosing variability in the Italian population. Pharmacogenomics, 2009, 10, 261-266.	0.6	129
2	Genetic Factors in Systemic Lupus Erythematosus: Contribution to Disease Phenotype. Journal of Immunology Research, 2015, 2015, 1-11.	0.9	79
3	Common polymorphisms in MIR146a, MIR128a and MIR27a genes contribute to neuropathy susceptibility in type 2 diabetes. Acta Diabetologica, 2014, 51, 663-671.	1.2	70
4	Nevirapine-induced hepatotoxicity and pharmacogenetics: a retrospective study in a population from Mozambique. Pharmacogenomics, 2010, 11, 23-31.	0.6	67
5	TCF7L2 gene polymorphisms and type 2 diabetes: association with diabetic retinopathy and cardiovascular autonomic neuropathy. Acta Diabetologica, 2013, 50, 789-799.	1.2	62
6	MicroRNA genetic variations: association with type 2 diabetes. Acta Diabetologica, 2013, 50, 867-872.	1.2	60
7	A Multilocus Genetic Study in a Cohort of Italian SLE Patients Confirms the Association with STAT4 Gene and Describes a New Association with HCP5 Gene. PLoS ONE, 2014, 9, e111991.	1.1	60
8	Allelic variants in theCYP2C9andVKORC1loci and interindividual variability in the anticoagulant dose effect of warfarin in Italians. Pharmacogenomics, 2007, 8, 1545-1550.	0.6	59
9	Impact of the CYP4F2 p.V433M Polymorphism on Coumarin Dose Requirement: Systematic Review and Meta-Analysis. Clinical Pharmacology and Therapeutics, 2012, 92, 746-756.	2.3	56
10	Association between CYP2B6 polymorphisms and Nevirapine-induced SJS/TEN: a pharmacogenetics study. European Journal of Clinical Pharmacology, 2013, 69, 1909-1916.	0.8	55
11	TRAF3IP2 gene and systemic lupus erythematosus: association with disease susceptibility and pericarditis development. Immunogenetics, 2013, 65, 703-709.	1.2	53
12	TRAF3IP2 gene is associated with cutaneous extraintestinal manifestations in Inflammatory Bowel Disease. Journal of Crohn's and Colitis, 2013, 7, 44-52.	0.6	51
13	Genetics and Treatment Response in Parkinsonâ€™s Disease: An Update on Pharmacogenetic Studies. NeuroMolecular Medicine, 2018, 20, 1-17.	1.8	43
14	Interaction between microbiome and host genetics in psoriatic arthritis. Autoimmunity Reviews, 2018, 17, 276-283.	2.5	38
15	Genetic tests and genomic biomarkers: regulation, qualification and validation. Clinical Cases in Mineral and Bone Metabolism, 2008, 5, 149-54.	1.0	37
16	Polymorphisms in STAT-4, IL-10, PSORS1C1, PTPN2 and MIR146A genes are associated differently with prognostic factors in Italian patients affected by rheumatoid arthritis. Clinical and Experimental Immunology, 2016, 186, 157-163.	1.1	36
17	Autophagy and inflammatory bowel disease: Association between variants of the autophagy-related IRGM gene and susceptibility to Crohn's disease. Digestive and Liver Disease, 2015, 47, 744-750.	0.4	35
18	Association between a MIR499A polymorphism and diabetic neuropathy in type 2 diabetes. Journal of Diabetes and Its Complications, 2018, 32, 11-17.	1.2	35

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19	Uncovering genetic and non-genetic biomarkers specific for exudative age-related macular degeneration: significant association of twelve variants. Oncotarget, 2018, 9, 7812-7821.	0.8	33
20	TNFAIP3 Gene Polymorphisms in Three Common Autoimmune Diseases: Systemic Lupus Erythematosus, Rheumatoid Arthritis, and Primary Sjogren Syndromeâ€"Association with Disease Susceptibility and Clinical Phenotypes in Italian Patients. Journal of Immunology Research, 2019, 2019, 1-6.	0.9	30
21	Recent advances in exploring the genetic susceptibility to diabetic neuropathy. Diabetes Research and Clinical Practice, 2016, 120, 198-208.	1.1	28
22	ATG16L1 Ala197Thr Is Not Associated With Susceptibility to Crohnâ€™s Disease or With Phenotype in an Italian Population. Gastroenterology, 2008, 134, 368-370.	0.6	26
23	Evaluation of ATG5 polymorphisms in Italian patients with systemic lupus erythematosus: contribution to disease susceptibility and clinical phenotypes. Lupus, 2018, 27, 1464-1469.	0.8	25
24	STAT4, TRAF3IP2, IL10, and HCP5 Polymorphisms in SjÃ¶grenâ€™s Syndrome: Association with Disease Susceptibility and Clinical Aspects. Journal of Immunology Research, 2019, 2019, 1-8.	0.9	25
25	Effect of <scp>CYP</scp>4F2,<scp>VKORC</scp>1, and <scp>CYP</scp>2C9 in Influencing Coumarin Dose: A Singleâ€Patient Data Metaâ€Analysis in More Than 15,000 Individuals. Clinical Pharmacology and Therapeutics, 2019, 105, 1477-1491.	2.3	23
26	Polymorphisms in miRNA genes and their involvement in autoimmune diseases susceptibility. Immunologic Research, 2017, 65, 811-827.	1.3	23
27	Polymorphisms in STAT4, PTPN2, PSORS1C1 and TRAF3IP2 Genes Are Associated with the Response to TNF Inhibitors in Patients with Rheumatoid Arthritis. PLoS ONE, 2017, 12, e0169956.	1.1	22
28	Interleukin-23R Arg381Gln Is Associated With Susceptibility to Crohnâ€™s Disease But Not With Phenotype in an Italian Population. Gastroenterology, 2007, 133, 1049-1051.	0.6	21
29	Expression study of candidate miRNAs and evaluation of their potential use as biomarkers of diabetic neuropathy. Epigenomics, 2020, 12, 575-585.	1.0	21
30	Haplotype block structure study of the CFTR gene. Most variants are associated with the M470 allele in several European populations. European Journal of Human Genetics, 2006, 14, 85-93.	1.4	20
31	Could MicroRNA polymorphisms influence warfarin dosing? A pharmacogenetics study on mir133 genes. Thrombosis Research, 2015, 136, 367-370.	0.8	20
32	miRNAs in drug response variability: potential utility as biomarkers for personalized medicine. Pharmacogenomics, 2019, 20, 1049-1059.	0.6	20
33	HCP5 genetic variant (RS3099844) contributes to Nevirapine-induced Stevens Johnsons Syndrome/Toxic Epidermal Necrolysis susceptibility in a population from Mozambique. European Journal of Clinical Pharmacology, 2014, 70, 275-278.	0.8	18
34	Mitochondrial DNA Copy Number in Peripheral Blood Is Reduced in Type 2 Diabetes Patients with Polyneuropathy and Associated with a MIR499A Gene Polymorphism. DNA and Cell Biology, 2020, 39, 1467-1472.	0.9	18
35	A large-scale study of the random variability of a coding sequence: a study on the CFTR gene. European Journal of Human Genetics, 2005, 13, 184-192.	1.4	17
36	Cannabinoid Poisoning by Hemp Seed Oil in a Child. Pediatric Emergency Care, 2017, 33, 344-345.	0.5	17

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37	Polymorphisms in MIR122, MIR196A2, and MIR124A Genes are Associated with Clinical Phenotypes in Inflammatory Bowel Diseases. Molecular Diagnosis and Therapy, 2017, 21, 107-114.	1.6	17
38	Highly preferential association of NonF508del CF mutations with the M470 allele. Journal of Cystic Fibrosis, 2007, 6, 15-22.	0.3	15
39	Characterization of a novel CYP2C9 gene mutation and structural bioinformatic protein analysis in a warfarin hypersensitive patient. Pharmacogenetics and Genomics, 2011, 21, 344-346.	0.7	15
40	Genetics, Epigenetics, and Gender Impact in Axial-Spondyloarthritis Susceptibility: An Update on Genetic Polymorphisms and Their Sex Related Associations. Frontiers in Genetics, 2021, 12, 671976.	1.1	14
41	A polymorphism upstream MIR1279 gene is associated with pericarditis development in Systemic Lupus Erythematosus and contributes to definition of a genetic risk profile for this complication. Lupus, 2017, 26, 841-848.	0.8	13
42	Impact of glutathione transferases genes polymorphisms in nevirapine adverse reactions: a possible role for GSTM1 in SJS/TEN susceptibility. European Journal of Clinical Pharmacology, 2017, 73, 1253-1259.	0.8	12
43	Pharmacogenetics of inflammatory bowel disease: a focus on Crohn's disease. Pharmacogenomics, 2017, 18, 1095-1114.	0.6	11
44	Pharmacogenomics: Role in Medicines Approval and Clinical Use. Public Health Genomics, 2010, 13, 284-291.	0.6	10
45	Stevensâ€“Johnson syndrome and toxic epidermal necrolysis: an update on pharmacogenetics studies in drug-induced severe skin reaction. Pharmacogenomics, 2015, 16, 1989-2002.	0.6	10
46	VDR Polymorphisms in Autoimmune Connective Tissue Diseases: Focus on Italian Population. Journal of Immunology Research, 2021, 2021, 1-6.	0.9	10
47	EPHX1 Polymorphisms Are Not Associated With Warfarin Response in an Italian Population. Clinical Pharmacology and Therapeutics, 2011, 89, 791-791.	2.3	9
48	Resequencing of VKORC1, CYP2C9 and CYP4F2 genes in Italian patients requiring extreme low and high warfarin doses. Thrombosis Research, 2013, 132, 123-126.	0.8	9
49	Emerging Role of microRNAs and Long Non-Coding RNAs in SjÃ¶grenâ€™s Syndrome. Genes, 2021, 12, 903.	1.0	9
50	Altered expression of miR-142, miR-155, miR-499a and of their putative common target MDM2 in systemic lupus erythematosus. Epigenomics, 2021, 13, 5-13.	1.0	8
51	What Is in the Field for Genetics and Epigenetics of Diabetic Neuropathy: The Role of MicroRNAs. Journal of Diabetes Research, 2021, 2021, 1-10.	1.0	8
52	Population differences in allele frequencies at theOLR1locus may suggest geographic disparities in cardiovascular risk events. Annals of Human Biology, 2010, 37, 137-149.	0.4	7
53	A Pharmacogenetics Study in Mozambican Patients Treated with Nevirapine: Full Resequencing of TRAF3IP2 Gene Shows a Novel Association with SJS/TEN Susceptibility. International Journal of Molecular Sciences, 2015, 16, 5830-5838.	1.8	7
54	A common polymorphism in MIR155 gene promoter region is associated with a lower risk to develop type 2 diabetes. Acta Diabetologica, 2019, 56, 717-718.	1.2	7

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55	The differential response to anti IL-6 treatment in COVID-19: the genetic counterpart. Clinical and Experimental Rheumatology, 2020, 38, 580.	0.4	7
56	Anthropological features of the CFTR gene: Its variability in an African population. Annals of Human Biology, 2011, 38, 203-209.	0.4	5
57	Advances in Exploring the Role of Micrornas in Inflammatory Bowel Disease. MicroRNA (Shariqah,) Tj ETQq1 1 0.784314 rgBT /Overlock	0.6	5
58	CLEC4E (Mincle) genetic variation associates with pulmonary tuberculosis in Guinea-Bissau (West) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50	1.0	5
59	mRNA expression analysis confirms CD44 splicing impairment in systemic lupus erythematosus patients. Lupus, 2021, 30, 1086-1093.	0.8	5
60	Pharmacogenomics in Parkinsonâ€™s disease: which perspective for developing a personalized medicine?. Neural Regeneration Research, 2019, 14, 75.	1.6	5
61	A family study of asymptomatic small bowel Crohn's disease. Digestive and Liver Disease, 2014, 46, 276-278.	0.4	4
62	The Impacts of the Clinical and Genetic Factors on Chronic Damage in Caucasian Systemic Lupus Erythematosus Patients. Journal of Clinical Medicine, 2022, 11, 3368.	1.0	4
63	A multilocus genetic study evidences the association of autoimmune-related genes with Psoriatic Arthritis in Italian patients. Immunobiology, 2022, 227, 152232.	0.8	3
64	A multivariate genetic analysis confirms rs5010528 in the human leucocyte antigen-C locus as a significant contributor to Stevens-Johnson syndrome/toxic epidermal necrolysis susceptibility in a Mozambique HIV population treated with nevirapine. Journal of Antimicrobial Chemotherapy, 2018, 73, 2137-2140.	1.3	2
65	ABCC10 rs2125739 polymorphism and nevirapine-induced hepatotoxicity. Pharmacogenetics and Genomics, 2013, 23, 38-39.	0.7	1
66	Impact of TRAF3IP2, IL10 and HCP5 Genetic Polymorphisms in the Response to TNF-i Treatment in Patients with Psoriatic Arthritis. Journal of Personalized Medicine, 2022, 12, 1094.	1.1	1
67	Genetics and Autoimmunity. , 2019, , 93-104.		0
68	Discovering the genetic contribution to cardiovascular diseases in patients affected by autoimmune diseases. Annals of Translational Medicine, 2018, 6, S44-S44.	0.7	0