

Valeria Novelli

List of Publications by Citations

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

28
papers

2,060
citations

18
h-index

31
g-index

31
ext. papers

2,593
ext. citations

7.2
avg, IF

3.99
L-index

#	Paper	IF	Citations
28	Missense mutations in plakophilin-2 cause sodium current deficit and associate with a Brugada syndrome phenotype. <i>Circulation</i> , 2014 , 129, 1092-103	16.7	242
27	Association of the FOXO3A locus with extreme longevity in a southern Italian centenarian study. <i>Rejuvenation Research</i> , 2009 , 12, 95-104	2.6	240
26	Frontotemporal dementia and its subtypes: a genome-wide association study. <i>Lancet Neurology, The</i> , 2014 , 13, 686-99	24.1	207
25	Adaptation and validation of the ACMG/AMP variant classification framework for MYH7-associated inherited cardiomyopathies: recommendations by ClinGen/Inherited Cardiomyopathy Expert Panel. <i>Genetics in Medicine</i> , 2018 , 20, 351-359	8.1	173
24	Reappraisal of Reported Genes for Sudden Arrhythmic Death: Evidence-Based Evaluation of Gene Validity for Brugada Syndrome. <i>Circulation</i> , 2018 , 138, 1195-1205	16.7	158
23	Novel insight into the natural history of short QT syndrome. <i>Journal of the American College of Cardiology</i> , 2014 , 63, 1300-1308	15.1	147
22	Gene-Specific Therapy With Mexiletine Reduces Arrhythmic Events in Patients With Long QT Syndrome Type 3. <i>Journal of the American College of Cardiology</i> , 2016 , 67, 1053-1058	15.1	123
21	Arrhythmogenic Right Ventricular Cardiomyopathy: Clinical Course and Predictors of Arrhythmic Risk. <i>Journal of the American College of Cardiology</i> , 2016 , 68, 2540-2550	15.1	99
20	An International, Multicentered, Evidence-Based Reappraisal of Genes Reported to Cause Congenital Long QT Syndrome. <i>Circulation</i> , 2020 , 141, 418-428	16.7	95
19	Association between a genetic variant related to glutamic acid metabolism and coronary heart disease in individuals with type 2 diabetes. <i>JAMA - Journal of the American Medical Association</i> , 2013 , 310, 821-8	27.4	95
18	Fatty acid profile of erythrocyte membranes as possible biomarker of longevity. <i>Rejuvenation Research</i> , 2008 , 11, 63-72	2.6	77
17	Mutational spectrum of the oral-facial-digital type I syndrome: a study on a large collection of patients. <i>Human Mutation</i> , 2008 , 29, 1237-46	4.7	73
16	Association study on long-living individuals from Southern Italy identifies rs10491334 in the CAMKIV gene that regulates survival proteins. <i>Rejuvenation Research</i> , 2011 , 14, 283-91	2.6	68
15	Lack of replication of genetic associations with human longevity. <i>Biogerontology</i> , 2008 , 9, 85-92	4.5	64
14	Association of rs2200733 at 4q25 with atrial flutter/fibrillation diseases in an Italian population. <i>Heart</i> , 2008 , 94, 1394-6	5.1	45
13	A genome-wide screening and SNPs-to-genes approach to identify novel genetic risk factors associated with frontotemporal dementia. <i>Neurobiology of Aging</i> , 2015 , 36, 2904.e13-26	5.6	34
12	Fatty acid percentage in erythrocyte membranes of atrial flutter/fibrillation patients and controls. <i>Journal of Interventional Cardiac Electrophysiology</i> , 2010 , 27, 95-9	2.4	29

11	A C6orf10/LOC101929163 locus is associated with age of onset in C9orf72 carriers. <i>Brain</i> , 2018 , 141, 2895-2907	11.2	25
10	Genetic modulators of the phenotype in the long QT syndrome: state of the art and clinical impact. <i>Current Opinion in Genetics and Development</i> , 2015 , 33, 17-24	4.9	17
9	A novel network analysis approach reveals DNA damage, oxidative stress and calcium/cAMP homeostasis-associated biomarkers in frontotemporal dementia. <i>PLoS ONE</i> , 2017 , 12, e0185797	3.7	14
8	Pleiotropic Phenotypes Associated With PKP2 Variants. <i>Frontiers in Cardiovascular Medicine</i> , 2018 , 5, 184	5.4	10
7	Role of extensive diagnostic workup in young athletes and nonathletes with complex ventricular arrhythmias. <i>Heart Rhythm</i> , 2020 , 17, 230-237	6.7	7
6	, age at onset, and ancestry help discriminate behavioral from language variants in FTLD cohorts. <i>Neurology</i> , 2020 , 95, e3288-e3302	6.5	5
5	Evaluation of gene validity for CPVT and short QT syndrome in sudden arrhythmic death. <i>European Heart Journal</i> , 2021 ,	9.5	5
4	Challenges in Molecular Diagnostics of Channelopathies in the Next-Generation Sequencing Era: Less Is More?. <i>Frontiers in Cardiovascular Medicine</i> , 2016 , 3, 29	5.4	4
3	Gene Expression Imputation Across Multiple Tissue Types Provides Insight Into the Genetic Architecture of Frontotemporal Dementia and Its Clinical Subtypes. <i>Biological Psychiatry</i> , 2021 , 89, 825-833	7.9	3
2	Mendelian randomization implies no direct causal association between leukocyte telomere length and amyotrophic lateral sclerosis. <i>Scientific Reports</i> , 2020 , 10, 12184	4.9	1
1	Clinical utility of genetic testing in the early diagnosis of Danon disease mimicking hypertrophic cardiomyopathy: a case report. <i>BMC Cardiovascular Disorders</i> , 2020 , 20, 156	2.3	