Valeria Novelli

List of Publications by Year
in descending order

31 papers	2,978 citations	361045 20 h-index	476904 29 g-index
31 all docs	31 docs citations	31 times ranked	6104 citing authors

#	Article	IF	Citations
1	Missense Mutations in Plakophilin-2 Cause Sodium Current Deficit and Associate With a Brugada Syndrome Phenotype. Circulation, 2014, 129, 1092-1103.	1.6	305
2	Frontotemporal dementia and its subtypes: a genome-wide association study. Lancet Neurology, The, 2014, 13, 686-699.	4.9	302
3	Adaptation and validation of the ACMG/AMP variant classification framework for MYH7-associated inherited cardiomyopathies: recommendations by ClinGenâ€™s Inherited Cardiomyopathy Expert Panel. Genetics in Medicine, 2018, 20, 351-359.	1.1	283
4	Association of theFOXO3ALocus with Extreme Longevity in a Southern Italian Centenarian Study. Rejuvenation Research, 2009, 12, 95-104.	0.9	282
5	Reappraisal of Reported Genes for Sudden Arrhythmic Death. Circulation, 2018, 138, 1195-1205.	1.6	271
6	An International, Multicentered, Evidence-Based Reappraisal of Genes Reported to Cause Congenital Long QT Syndrome. Circulation, 2020, 141, 418-428.	1.6	238
7	Novel Insight Into the Natural History of Short QT Syndrome. Journal of the American College of Cardiology, 2014, 63, 1300-1308.	1.2	191
8	Gene-Specific Therapy With Mexiletine Reduces Arrhythmic Events in Patients With Long QT Syndrome Type 3. Journal of the American College of Cardiology, 2016, 67, 1053-1058.	1.2	191
9	Arrhythmogenic Right Ventricular Cardiomyopathy. Journal of the American College of Cardiology, 2016, 68, 2540-2550.	1.2	148
10	Association Between a Genetic Variant Related to Glutamic Acid Metabolism and Coronary Heart Disease in Individuals With Type 2 Diabetes. JAMA - Journal of the American Medical Association, 2013, 310, 821.	3.8	122
11	Fatty Acid Profile of Erythrocyte Membranes As Possible Biomarker of Longevity. Rejuvenation Research, 2008, 11, 63-72.	0.9	87
12	Mutational spectrum of the oral-facial-digital type I syndrome: a study on a large collection of patients. Human Mutation, 2008, 29, 1237-1246.	1.1	82
13	Association Study on Long-Living Individuals from Southern Italy Identifies rs10491334 in the CAMKIV Gene That Regulates Survival Proteins. Rejuvenation Research, 2011, 14, 283-291.	0.9	72
14	Lack of replication of genetic associations with human longevity. Biogerontology, 2008, 9, 85-92.	2.0	69
15	Evaluation of gene validity for CPVT and short QT syndrome in sudden arrhythmic death. European Heart Journal, 2022, 43, 1500-1510.	1.0	57
16	Association of rs2200733 at 4q25 with atrial flutter/fibrillation diseases in an Italian population. Heart, 2008, 94, 1394-1396.	1.2	49
17	A genome-wide screening and SNPs-to-genes approach to identify novel genetic risk factors associated with frontotemporal dementia. Neurobiology of Aging, 2015, 36, 2904.e13-2904.e26.	1.5	48
18	A C6orf10/LOC101929163 locus is associated with age of onset in C9orf72 carriers. Brain, 2018, 141, 2895-2907.	3.7	39

#	Article	IF	Citations
19	Fatty acid percentage in erythrocyte membranes of atrial flutter/fibrillation patients and controls. Journal of Interventional Cardiac Electrophysiology, 2010, 27, 95-99.	0.6	32
20	Pleiotropic Phenotypes Associated With PKP2 Variants. Frontiers in Cardiovascular Medicine, 2018, 5, 184.	1.1	23
21	A novel network analysis approach reveals DNA damage, oxidative stress and calcium/cAMP homeostasis-associated biomarkers in frontotemporal dementia. PLoS ONE, 2017, 12, e0185797.	1.1	21
22	Genetic modulators of the phenotype in the long QT syndrome: state of the art and clinical impact. Current Opinion in Genetics and Development, 2015, 33, 17-24.	1.5	19
23	Role of extensive diagnostic workup in young athletes and nonathletes with complex ventricular arrhythmias. Heart Rhythm, 2020, 17, 230-237.	0.3	10
24	Gene Expression Imputation Across Multiple Tissue Types Provides Insight Into the Genetic Architecture of Frontotemporal Dementia and Its Clinical Subtypes. Biological Psychiatry, 2021, 89, 825-835.	0.7	10
25	Challenges in Molecular Diagnostics of Channelopathies in the Next-Generation Sequencing Era: Less Is More?. Frontiers in Cardiovascular Medicine, 2016, 3, 29.	1.1	8
26	C9orf72, age at onset, and ancestry help discriminate behavioral from language variants in FTLD cohorts. Neurology, 2020, 95, e3288-e3302.	1.5	7
27	Role of CACNA1C in Brugada syndrome: Prevalence and phenotype of probands referred for genetic testing. Heart Rhythm, 2022, 19, 798-806.	0.3	7
28	Mendelian randomization implies no direct causal association between leukocyte telomere length and amyotrophic lateral sclerosis. Scientific Reports, 2020, 10, 12184.	1.6	4
29	Editorial: Current Challenges in Cardiovascular Molecular Diagnostics. Frontiers in Cardiovascular Medicine, 2017, 4, 54.	1.1	1
30	Clinical utility of genetic testing in the early diagnosis of Danon disease mimicking hypertrophic cardiomyopathy: a case report. BMC Cardiovascular Disorders, 2020, 20, 156.	0.7	0
31	Diagnostic Workflow in Competitive Athletes with Ventricular Arrhythmias and Suspected Concealed Cardiomyopathies. Medicina (Lithuania), 2021, 57, 182.	0.8	0