Irene Mademont

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/5507223/publications.pdf

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16	309	932766 10	996533 15
papers	citations	h-index	g-index

16	16	16	579
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Natural and Undetermined Sudden Death: Value of Post-Mortem Genetic Investigation. PLoS ONE, 2016, 11, e0167358.	1.1	62
2	Post-mortem genetic analysis in juvenile cases of sudden cardiac death. Forensic Science International, 2014, 245, 30-37.	1.3	44
3	Additional value of screening for minor genes and copy number variants in hypertrophic cardiomyopathy. PLoS ONE, 2017, 12, e0181465.	1.1	32
4	Identification of Genetic Alterations, as Causative Genetic Defects in Long QT Syndrome, Using Next Generation Sequencing Technology. PLoS ONE, 2014, 9, e114894.	1.1	26
5	Role of copy number variants in sudden cardiac death and related diseases: genetic analysis and translation into clinical practice. European Journal of Human Genetics, 2018, 26, 1014-1025.	1.4	26
6	Comprehensive Genetic Characterization of a Spanish Brugada Syndrome Cohort. PLoS ONE, 2015, 10, e0132888.	1.1	25
7	Large Genomic Imbalances in Brugada Syndrome. PLoS ONE, 2016, 11, e0163514.	1.1	23
8	Sudden Cardiac Death and Copy Number Variants: What Do We Know after 10 Years of Genetic Analysis?. Forensic Science International: Genetics, 2020, 47, 102281.	1.6	20
9	Rare Titin (TTN) Variants in Diseases Associated with Sudden Cardiac Death. International Journal of Molecular Sciences, 2015, 16, 25773-25787.	1.8	16
10	Sudden Arrhythmic Death During Exercise: A Post-Mortem Genetic Analysis. Sports Medicine, 2017, 47, 2101-2115.	3.1	11
11	Molecular autopsy in a cohort of infants died suddenly at rest. Forensic Science International: Genetics, 2018, 37, 54-63.	1.6	10
12	Personalized Interpretation and Clinical Translation of Genetic Variants Associated With Cardiomyopathies. Frontiers in Genetics, 2019, 10, 450.	1.1	6
13	GLYT1 encephalopathy: Further delineation of disease phenotype and discussion of pathophysiological mechanisms. American Journal of Medical Genetics, Part A, 2021, 185, 476-485.	0.7	5
14	A novel variant in RyR2 causes familiar catecholaminergic polymorphic ventricular tachycardia. Forensic Science International, 2017, 270, 173-177.	1.3	2
15	Genetic analysis in post-mortem samples with micro-ischemic alterations. Forensic Science International, 2017, 271, 120-125.	1.3	1
16	GRAPES: A Versatile Tool for Analyzing Structural Variation From Wholeâ€Genome and Targeted DNA Sequencing Data. FASEB Journal, 2018, 32, 532.10.	0.2	0