

Maria Gnoli

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5223384/publications.pdf>

Version: 2024-02-01

11
papers

116
citations

1478505

6
h-index

1474206

9
g-index

11
all docs

11
docs citations

11
times ranked

190
citing authors

#	ARTICLE	IF	CITATIONS
1	Genotype-phenotype correlation study in 364 osteogenesis imperfecta Italian patients. <i>European Journal of Human Genetics</i> , 2019, 27, 1090-1100.	2.8	52
2	Patterns of Novel Alleles and Genotype/Phenotype Correlations Resulting from the Analysis of 108 Previously Undetected Mutations in Patients Affected by Neurofibromatosis Type I. <i>International Journal of Molecular Sciences</i> , 2017, 18, 2071.	4.1	11
3	The natural history of multiple osteochondromas in a large Italian cohort of pediatric patients. <i>Bone</i> , 2020, 139, 115499.	2.9	11
4	COL1-Related Disorders: Case Report and Review of Overlapping Syndromes. <i>Frontiers in Genetics</i> , 2021, 12, 640558.	2.3	9
5	Melorheostosis and Osteopoikilosis Clinical and Molecular Description of an Italian Case Series. <i>Calcified Tissue International</i> , 2019, 105, 215-221.	3.1	7
6	Variable clinical expression of Stickler Syndrome: A case report of a novel <i>COL11A1</i> mutation. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1353.	1.2	7
7	Tumor Syndromes That Include Bone Tumors. <i>Surgical Pathology Clinics</i> , 2017, 10, 749-764.	1.7	6
8	The Rizzoli Multiple Osteochondromas Classification revised: describing the phenotype to improve clinical practice. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 3466-3475.	1.2	6
9	Evaluation of TP53 Pro72Arg and MDM2 SNP285-SNP309 polymorphisms in an Italian cohort of LFS suggestive patients lacking identifiable TP53 germline mutations. <i>Familial Cancer</i> , 2016, 15, 635-643.	1.9	4
10	Familial osteopoikilosis: Case report with differential diagnosis and review of the literature. <i>Clinical Case Reports (discontinued)</i> , 2021, 9, 922-926.	0.5	2
11	An Easy-to-Use Approach to Detect CNV From Targeted NGS Data: Identification of a Novel Pathogenic Variant in MO Disease. <i>Frontiers in Endocrinology</i> , 0, 13, .	3.5	1