

Benoit Arveiler

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

1,339
citations

471509

17
h-index

395702

33
g-index

37
all docs

37
docs citations

37
times ranked

2638
citing authors

#	ARTICLE	IF	CITATIONS
1	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. <i>Nature</i> , 2011, 478, 97-102.	27.8	394
2	Increasing the complexity: new genes and new types of albinism. <i>Pigment Cell and Melanoma Research</i> , 2014, 27, 11-18.	3.3	179
3	Molecular characterization of a series of 990 index patients with albinism. <i>Pigment Cell and Melanoma Research</i> , 2018, 31, 466-474.	3.3	92
4	Oculocutaneous albinism with TYRP1 gene mutations in a Caucasian patient. <i>Pigment Cell & Melanoma Research</i> , 2006, 19, 239-242.	3.6	62
5	Array-CGH analysis of a cohort of 86 patients with oculoauriculovertrebral spectrum. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 1984-1989.	1.2	53
6	Mutations in <i>MYT1</i> , encoding the myelin transcription factor 1, are a rare cause of OAVS. <i>Journal of Medical Genetics</i> , 2016, 53, 752-760.	3.2	51
7	New candidate loci identified by array-CGH in a cohort of 100 children presenting with syndromic obesity. <i>American Journal of Medical Genetics, Part A</i> , 2014, 164, 1965-1975.	1.2	49
8	BLOC1S5 pathogenic variants cause a new type of Hermansky-Pudlak syndrome. <i>Genetics in Medicine</i> , 2020, 22, 1613-1622.	2.4	44
9	Spectrum of CREBBP gene dosage anomalies in Rubinstein-Taybi Syndrome patients. <i>European Journal of Human Genetics</i> , 2007, 15, 843-847.	2.8	41
10	SLC24A5 Mutations Are Associated with Non-Syndromic Oculocutaneous Albinism. <i>Journal of Investigative Dermatology</i> , 2014, 134, 568-571.	0.7	36
11	A novel de novo mutation in MYT1, the unique OAVS gene identified so far. <i>European Journal of Human Genetics</i> , 2017, 25, 1083-1086.	2.8	35
12	Dopachrome tautomerase variants in patients with oculocutaneous albinism. <i>Genetics in Medicine</i> , 2021, 23, 479-487.	2.4	33
13	Complete loss of function of the ubiquitin ligase HERC2 causes a severe neurodevelopmental phenotype. <i>European Journal of Human Genetics</i> , 2017, 25, 52-58.	2.8	28
14	Mild form of oculocutaneous albinism type 1: phenotypic analysis of compound heterozygous patients with the R402Q variant of the <i>TYR</i> gene. <i>British Journal of Ophthalmology</i> , 2019, 103, 1239-1247.	3.9	24
15	High-resolution array-CGH in patients with oculocutaneous albinism identifies new deletions of the <i>TYR</i> , <i>OCA2</i> , and <i>SLC45A2</i> genes and a complex rearrangement of the <i>OCA2</i> gene. <i>Pigment Cell and Melanoma Research</i> , 2014, 27, 59-71.	3.3	23
16	Functional and genetic analyses of <i>ZYG11B</i> provide evidences for its involvement in OAVS. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1375.	1.2	21
17	Identification of a homozygous mutation of SLC24A5 (<i>OCA6</i>) in two patients with oculocutaneous albinism from French Guiana. <i>Pigment Cell and Melanoma Research</i> , 2016, 29, 104-106.	3.3	19
18	The contribution of common regulatory and protein-coding TYR variants to the genetic architecture of albinism. <i>Nature Communications</i> , 2022, 13, .	12.8	17

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19	High resolution mapping of OCA2 intragenic rearrangements and identification of a founder effect associated with a deletion in Polish albino patients. <i>Human Genetics</i> , 2011, 129, 199-208.	3.8	14
20	A recurrent missense variant in EYA3 gene is associated with oculo-auriculo-vertebral spectrum. <i>Human Genetics</i> , 2021, 140, 933-944.	3.8	14
21	Expanding the clinical phenotype at the 3q13.31 locus with a new case of microdeletion and first characterization of the reciprocal duplication. <i>Molecular Genetics and Metabolism</i> , 2013, 110, 90-97.	1.1	13
22	Clinico-molecular analysis of eleven patients with Hermansky-Pudlak type 5 syndrome, a mild form of <scp>HPS</scp>. <i>Pigment Cell and Melanoma Research</i> , 2017, 30, 563-570.	3.3	13
23	Hereditary Mucoepithelial Dysplasia Results from Heterozygous Variants at p.Arg557 Mutational Hotspot in SREBF1, Encoding a Transcription Factor Involved in Cholesterol Homeostasis. <i>Journal of Investigative Dermatology</i> , 2020, 140, 1289-1292.e2.	0.7	12
24	The Phenotypic and Mutational Spectrum of the FHONDA Syndrome and Oculocutaneous Albinism: Similarities and Differences. , 2022, 63, 19.		12
25	Albinism: An Underdiagnosed Condition. <i>Journal of Investigative Dermatology</i> , 2020, 140, 1449-1451.	0.7	9
26	19p13 microduplications encompassing NFIX are responsible for intellectual disability, short stature and small head circumference. <i>European Journal of Human Genetics</i> , 2018, 26, 85-93.	2.8	7
27	A new case with Hermansky-Pudlak syndrome type 9, a rare cause of syndromic albinism with severe defect of platelets dense bodies. <i>Platelets</i> , 2021, 32, 420-423.	2.3	5
28	Novel variants in the <i>BLOC1S3</i> gene in patients presenting a mild form of Hermansky-Pudlak syndrome. <i>Pigment Cell and Melanoma Research</i> , 2021, 34, 132-135.	3.3	5
29	Albinism in a patient with mutations at both the OA1 and OCA3 loci. <i>Pigment Cell and Melanoma Research</i> , 2016, 29, 107-109.	3.3	3
30	Deletion in 2q35 excluding the IHH gene leads to fetal severe limb anomalies and suggests a disruption of chromatin architecture. <i>European Journal of Human Genetics</i> , 2019, 27, 384-388.	2.8	3
31	The Dct ^{+/+} Mouse Model to Unravel Retinogenesis Misregulation in Patients with Albinism. <i>Genes</i> , 2022, 13, 1164.	2.4	3
32	Clinical and molecular findings of FRMD7 related congenital nystagmus as a differential diagnosis of ocular albinism. <i>Ophthalmic Genetics</i> , 2019, 40, 161-164.	1.2	2
33	Albinism. , 2022, , 393-402.		2
34	Clinical variability and probable founder effect in oculocutaneous albinism type 7. <i>Clinical Genetics</i> , 2020, 97, 527-528.	2.0	1