

Anne-Louise Leutenegger

List of Publications by Year in descending order

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Version: 2024-02-01

45
papers

3,845
citations

236833

25
h-index

206029

48
g-index

50
all docs

50
docs citations

50
times ranked

7447
citing authors

#	ARTICLE	IF	CITATIONS
1	Multiethnic genome-wide association study of differentiated thyroid cancer in the EPITHYR consortium. <i>International Journal of Cancer</i> , 2021, 148, 2935-2946.	2.3	11
2	Genetics of PlGF plasma levels highlights a role of its receptors and supports the link between angiogenesis and immunity. <i>Scientific Reports</i> , 2021, 11, 16821.	1.6	6
3	Clinical interpretation of variants identified in RNU4ATAC, a non-coding spliceosomal gene. <i>PLoS ONE</i> , 2020, 15, e0235655.	1.1	8
4	Novel Chronic Mouse Model of Cerebral Cavernous Malformations. <i>Stroke</i> , 2020, 51, 1272-1278.	1.0	25
5	New insights into minor splicing—a transcriptomic analysis of cells derived from TALS patients. <i>Rna</i> , 2019, 25, 1130-1149.	1.6	27
6	Whole-Exome Sequencing in the Isolated Populations of Cilento from South Italy. <i>Scientific Reports</i> , 2019, 9, 4059.	1.6	7
7	Strategies for phasing and imputation in a population isolate. <i>Genetic Epidemiology</i> , 2018, 42, 201-213.	0.6	27
8	Detecting the dominance component of heritability in isolated and outbred human populations. <i>Scientific Reports</i> , 2018, 8, 18048.	1.6	3
9	Variation in worldwide incidence of amyotrophic lateral sclerosis: a meta-analysis. <i>International Journal of Epidemiology</i> , 2017, 46, dyw061.	0.9	202
10	Loss of VPS13C Function in Autosomal-Recessive Parkinsonism Causes Mitochondrial Dysfunction and Increases PINK1/Parkin-Dependent Mitophagy. <i>American Journal of Human Genetics</i> , 2016, 98, 500-513.	2.6	333
11	Relationship inference from the genetic data on parents or offspring: A comparative study. <i>Theoretical Population Biology</i> , 2016, 107, 31-38.	0.5	1
12	Clinical and demographic factors and outcome of amyotrophic lateral sclerosis in relation to population ancestral origin. <i>European Journal of Epidemiology</i> , 2016, 31, 229-245.	2.5	87
13	Six Novel Loci Associated with Circulating VEGF Levels Identified by a Meta-analysis of Genome-Wide Association Studies. <i>PLoS Genetics</i> , 2016, 12, e1005874.	1.5	56
14	High level of inbreeding in final phase of 1000 Genomes Project. <i>Scientific Reports</i> , 2015, 5, 17453.	1.6	68
15	Genome-wide inbreeding estimation within Lebanese communities using SNP arrays. <i>European Journal of Human Genetics</i> , 2015, 23, 1364-1369.	1.4	11
16	Genetic Variants Modulating CRIPTO Serum Levels Identified by Genome-Wide Association Study in Cilento Isolates. <i>PLoS Genetics</i> , 2015, 11, e1004976.	1.5	13
17	A new F-box protein 7 gene mutation causing typical Parkinson's disease. <i>Movement Disorders</i> , 2015, 30, 1130-1133.	2.2	59
18	FSuite: exploiting inbreeding in dense SNP chip and exome data. <i>Bioinformatics</i> , 2014, 30, 1940-1941.	1.8	30

#	ARTICLE	IF	CITATIONS
19	Does anonymous sperm donation increase the risk for unions between relatives and the incidence of autosomal recessive diseases due to consanguinity?. <i>Human Reproduction</i> , 2014, 29, 394-399.	0.4	7
20	Inbreeding Coefficient Estimation with Dense SNP Data: Comparison of Strategies and Application to HapMap III. <i>Human Heredity</i> , 2014, 77, 49-62.	0.4	46
21	Could Inbred Cases Identified in GWAS Data Succeed in Detecting Rare Recessive Variants Where Affected Sib-Pairs Have Failed?. <i>Human Heredity</i> , 2012, 74, 142-152.	0.4	8
22	Genetic and Environmental Factors Influencing the Placental Growth Factor (PGF) Variation in Two Populations. <i>PLoS ONE</i> , 2012, 7, e42537.	1.1	11
23	Genetics of VEGF Serum Variation in Human Isolated Populations of Cilento: Importance of VEGF Polymorphisms. <i>PLoS ONE</i> , 2011, 6, e16982.	1.1	68
24	Consanguinity around the world: what do the genomic data of the HGDP-CEPH diversity panel tell us?. <i>European Journal of Human Genetics</i> , 2011, 19, 583-587.	1.4	52
25	Does inbreeding affect N-glycosylation of human plasma proteins?. <i>Molecular Genetics and Genomics</i> , 2011, 285, 427-432.	1.0	2
26	Polynesian ecology determines seasonality of biliary atresia. <i>Hepatology</i> , 2011, 54, 1893-1894.	3.6	12
27	Association of TALS Developmental Disorder with Defect in Minor Splicing Component <i>U4atac</i> snRNA. <i>Science</i> , 2011, 332, 240-243.	6.0	195
28	Comparative assessment of methods for estimating individual genome-wide homozygosity-by-descent from human genomic data. <i>BMC Genomics</i> , 2010, 11, 139.	1.2	29
29	Parkinson's disease-related LRRK2 G2019S mutation results from independent mutational events in humans. <i>Human Molecular Genetics</i> , 2010, 19, 1998-2004.	1.4	48
30	Runs of Homozygosity in European Populations. <i>American Journal of Human Genetics</i> , 2008, 83, 359-372.	2.6	958
31	Runs of Homozygosity in European Populations. <i>American Journal of Human Genetics</i> , 2008, 83, 658.	2.6	13
32	LRRK2emph Exon 41 Mutations in Sporadic Parkinson Disease in Europeans. <i>Archives of Neurology</i> , 2007, 64, 425.	4.9	51
33	A novel locus for autosomal dominant "uncomplicated" hereditary spastic paraplegia maps to chromosome 8p21.1-q13.3. <i>Human Genetics</i> , 2007, 122, 261-273.	1.8	27
34	Using Genomic Inbreeding Coefficient Estimates for Homozygosity Mapping of Rare Recessive Traits: Application to Taybi-Linder Syndrome. <i>American Journal of Human Genetics</i> , 2006, 79, 62-66.	2.6	48
35	LRRK2G2019S as a Cause of Parkinson's Disease in North African Arabs. <i>New England Journal of Medicine</i> , 2006, 354, 422-423.	13.9	521
36	Juvenile-Onset Parkinsonism as a Result of the First Mutation in the Adenosine Triphosphate Orientation Domain of PINK1. <i>Archives of Neurology</i> , 2006, 63, 1257.	4.9	43

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37	G2019S LRRK2 mutation in French and North African families with Parkinson's disease. <i>Annals of Neurology</i> , 2005, 58, 784-787.	2.8	196
38	Detection of susceptibility loci by genome-wide linkage analysis. <i>BMC Genetics</i> , 2005, 6, S18.	2.7	5
39	Modeling the effect of a genetic factor for a complex trait in a simulated population. <i>BMC Genetics</i> , 2005, 6, S87.	2.7	1
40	LRRK2 Haplotype Analyses in European and North African Families with Parkinson Disease: A Common Founder for the G2019S Mutation Dating from the 13th Century. <i>American Journal of Human Genetics</i> , 2005, 77, 330-332.	2.6	130
41	Estimation of the Inbreeding Coefficient through Use of Genomic Data. <i>American Journal of Human Genetics</i> , 2003, 73, 516-523.	2.6	221
42	Presence of Large Deletions in Kindreds with Autism. <i>American Journal of Human Genetics</i> , 2002, 71, 100-115.	2.6	63
43	Impact of parental relationships in maximum lod score affected sib-pair method. <i>Genetic Epidemiology</i> , 2002, 23, 413-425.	0.6	25
44	The Importance of Connections: Joining Components of the Hutterite Pedigree. <i>Genetic Epidemiology</i> , 2001, 21, S230-5.	0.6	11
45	Segregation Analysis of Phenotypic Components of Learning Disabilities. I. Nonword Memory and Digit Span. <i>American Journal of Human Genetics</i> , 2000, 67, 631-646.	2.6	73