

# Francesco Lescai

## List of Publications by Year in descending order

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Version: 2024-02-01

46  
papers

2,417  
citations

218662

26  
h-index

302107

39  
g-index

53  
all docs

53  
docs citations

53  
times ranked

6308  
citing authors

#	ARTICLE	IF	CITATIONS
1	Novel variation and de novo mutation rates in population-wide de novo assembled Danish trios. <i>Nature Communications</i> , 2015, 6, 5969.	12.8	164
2	Common variants at the MHC locus and at chromosome 16q24.1 predispose to Barrett's esophagus. <i>Nature Genetics</i> , 2012, 44, 1131-1136.	21.4	162
3	Autism spectrum disorder and attention deficit hyperactivity disorder have a similar burden of rare protein-truncating variants. <i>Nature Neuroscience</i> , 2019, 22, 1961-1965.	14.8	148
4	Human models of aging and longevity. <i>Expert Opinion on Biological Therapy</i> , 2008, 8, 1393-1405.	3.1	147
5	Experimental validation of methods for differential gene expression analysis and sample pooling in RNA-seq. <i>BMC Genomics</i> , 2015, 16, 548.	2.8	139
6	Do men and women follow different trajectories to reach extreme longevity?. <i>Aging Clinical and Experimental Research</i> , 2000, 12, 77-84.	2.9	138
7	Sequencing and de novo assembly of 150 genomes from Denmark as a population reference. <i>Nature</i> , 2017, 548, 87-91.	27.8	130
8	Association between the interleukin-1 $\beta$ polymorphisms and Alzheimer's disease: A systematic review and meta-analysis. <i>Brain Research Reviews</i> , 2008, 59, 155-163.	9.0	107
9	Identification of germline susceptibility loci in ETV6-RUNX1-rearranged childhood acute lymphoblastic leukemia. <i>Leukemia</i> , 2012, 26, 902-909.	7.2	106
10	Quantifying the Impact of Rare and Ultra-rare Coding Variation across the Phenotypic Spectrum. <i>American Journal of Human Genetics</i> , 2018, 102, 1204-1211.	6.2	102
11	Mutations in SNX14 Cause a Distinctive Autosomal-Recessive Cerebellar Ataxia and Intellectual Disability Syndrome. <i>American Journal of Human Genetics</i> , 2014, 95, 611-621.	6.2	89
12	Systems Biology and Longevity: An Emerging Approach to Identify Innovative Anti- Aging Targets and Strategies. <i>Current Pharmaceutical Design</i> , 2010, 16, 802-813.	1.9	76
13	The kinetochore protein, <i>CENPF</i> , is mutated in human ciliopathy and microcephaly phenotypes. <i>Journal of Medical Genetics</i> , 2015, 52, 147-156.	3.2	75
14	Complexity of Anti-immunosenesence Strategies in Humans. <i>Artificial Organs</i> , 2006, 30, 730-742.	1.9	68
15	Genes, ageing and longevity in humans: Problems, advantages and perspectives. <i>Free Radical Research</i> , 2006, 40, 1303-1323.	3.3	66
16	ARNT2 mutation causes hypopituitarism, post-natal microcephaly, visual and renal anomalies. <i>Brain</i> , 2013, 136, 3096-3105.	7.6	66
17	Human longevity and 11p15.5: a study in 1321 centenarians. <i>European Journal of Human Genetics</i> , 2009, 17, 1515-1519.	2.8	60
18	PON1 is a longevity gene: Results of a meta-analysis. <i>Ageing Research Reviews</i> , 2009, 8, 277-284.	10.9	59

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19	An APOE Haplotype Associated with Decreased $\beta$ 4 Expression Increases the Risk of Late Onset Alzheimer's Disease. <i>Journal of Alzheimer's Disease</i> , 2011, 24, 235-245.	2.6	58
20	STAG3 truncating variant as the cause of primary ovarian insufficiency. <i>European Journal of Human Genetics</i> , 2016, 24, 135-138.	2.8	53
21	Neuroinflammation and the genetics of Alzheimer's disease: The search for a pro-inflammatory phenotype. <i>Aging Clinical and Experimental Research</i> , 2001, 13, 163-170.	2.9	44
22	Mutation of SALL2 causes recessive ocular coloboma in humans and mice. <i>Human Molecular Genetics</i> , 2014, 23, 2511-2526.	2.9	39
23	High-Quality Exome Sequencing of Whole-Genome Amplified Neonatal Dried Blood Spot DNA. <i>PLoS ONE</i> , 2016, 11, e0153253.	2.5	38
24	Identification of single nucleotide polymorphisms in the p21 (CDKN1A) gene and correlations with longevity in the Italian population. <i>Aging</i> , 2009, 1, 470-480.	3.1	34
25	A Genetic-Demographic Approach Reveals Male-Specific Association Between Survival and Tumor Necrosis Factor (A/G)-308 Polymorphism. <i>Journals of Gerontology - Series A Biological Sciences and Medical Sciences</i> , 2008, 63, 454-460.	3.6	30
26	ParkDB: a Parkinson's disease gene expression database. <i>Database: the Journal of Biological Databases and Curation</i> , 2011, 2011, bar007-bar007.	3.0	28
27	Whole-Exome Sequencing Reveals Increased Burden of Rare Functional and Disruptive Variants in Candidate Risk Genes in Individuals With Persistent Attention-Deficit/Hyperactivity Disorder. <i>Journal of the American Academy of Child and Adolescent Psychiatry</i> , 2016, 55, 521-523.	0.5	28
28	Whole-exome sequencing of individuals from an isolated population implicates rare risk variants in bipolar disorder. <i>Translational Psychiatry</i> , 2017, 7, e1034-e1034.	4.8	24
29	The Impact of Phenocopy on the Genetic Analysis of Complex Traits. <i>PLoS ONE</i> , 2010, 5, e11876.	2.5	19
30	The use of whole-exome sequencing to disentangle complex phenotypes. <i>European Journal of Human Genetics</i> , 2016, 24, 298-301.	2.8	15
31	Assembly and analysis of 100 full MHC haplotypes from the Danish population. <i>Genome Research</i> , 2017, 27, 1597-1607.	5.5	15
32	Whole-exome sequencing implicates <i>DGKH</i> as a risk gene for panic disorder in the Faroese population. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 1013-1022.	1.7	14
33	Identification and validation of loss of function variants in clinical contexts. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2014, 2, 58-63.	1.2	12
34	Failure to Replicate an Association of rs5984894 SNP in the PCDH11X Gene in a Collection of 1,222 Alzheimer's Disease Affected Patients. <i>Journal of Alzheimer's Disease</i> , 2010, 21, 385-388.	2.6	11
35	Characterisation and Validation of Insertions and Deletions in 173 Patient Exomes. <i>PLoS ONE</i> , 2012, 7, e51292.	2.5	8
36	Paraoxonase-1 55 LL Genotype Is Associated with No ST-Elevation Myocardial Infarction and with High Levels of Myoglobin. <i>Journal of Lipids</i> , 2012, 2012, 1-5.	4.8	6

#	ARTICLE	IF	CITATIONS
37	Omics in a Digital World: The Role of Bioinformatics in Providing New Insights Into Human Aging. <i>Frontiers in Genetics</i> , 2021, 12, 689824.	2.3	6
38	Genotype of inflammatory cytokines in limbal stem cell graft in Italian patients. <i>Biochemical and Biophysical Research Communications</i> , 2005, 332, 95-100.	2.1	3
39	Mutations in SNX14 Cause a Distinctive Autosomal-Recessive Cerebellar Ataxia and Intellectual Disability Syndrome. <i>American Journal of Human Genetics</i> , 2015, 96, 1008-1009.	6.2	1
40	Aging and Longevity in Animal Models and Humans. , 2009, , 175-191.		1
41	Italian biotechnologists organize. <i>Nature</i> , 2003, 425, 644-644.	27.8	0
42	Pathological Relevance of the Natural Immune System. <i>NeuroImmune Biology</i> , 2005, , 331-350.	0.2	0
43	Helping young independent scientists: the EMBO Young Investigator Programme. <i>New Biotechnology</i> , 2008, 25, 120-121.	4.4	0
44	The Young European Biotech Network (YEBN). <i>New Biotechnology</i> , 2008, 25, 34.	4.4	0
45	Special issue of new biotechnology in memory of professor Brian F.C. Clark (1936â€“2014). <i>New Biotechnology</i> , 2017, 38, 1.	4.4	0
46	Role of mitochondrial DNA in longevity, aging and age-related diseases in humans: a reappraisal. <i>Italian Journal of Biochemistry</i> , 2007, 56, 243-53.	0.3	0