

# Luigi Bisceglia

## List of Publications by Year in descending order

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55  
papers

3,367  
citations

257450

24  
h-index

182427

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55  
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55  
docs citations

55  
times ranked

3618  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mitochondrial DNA copy number in affected and unaffected LHON mutation carriers. BMC Research Notes, 2018, 11, 911.	1.4	25
2	Leber's hereditary optic neuropathy, intellectual disability and epilepsy presenting with variable penetrance associated to the m.3460C>A mutation and a heteroplasmic expansion of the microsatellite in MTRNR1 gene – case report. BMC Medical Genetics, 2018, 19, 129.	2.1	7
3	Leber's hereditary optic neuropathy (LHON) in an Apulian cohort of subjects. Acta Myologica, 2017, 36, 163-177.	1.5	7
4	Expanding the mutation spectrum in 130 probands with ARPKD: identification of 62 novel PKHD1 mutations by sanger sequencing and MLPA analysis. Journal of Human Genetics, 2016, 61, 811-821.	2.3	27
5	Mitochondrial DNA copy number differentiates the Leber's hereditary optic neuropathy affected individuals from the unaffected mutation carriers. Brain, 2016, 139, e1-e1.	7.6	39
6	Identification and Clinical Characterization of Adult Patients with Multigenerational Diabetes Mellitus. PLoS ONE, 2015, 10, e0135855.	2.5	14
7	20 novel point mutations and one large deletion in EXT1 and EXT2 genes: Report of diagnostic screening in a large Italian cohort of patients affected by hereditary multiple exostosis. Gene, 2013, 515, 339-348.	2.2	24
8	Pontocerebellar hypoplasia type 6 caused by mutations in <i>RARS2</i> : definition of the clinical spectrum and molecular findings in five patients. Journal of Inherited Metabolic Disease, 2013, 36, 43-53.	3.6	70
9	Clinical utility gene card for: Cystinuria. European Journal of Human Genetics, 2012, 20, 3-3.	2.8	12
10	Clinical, biochemical and molecular characterization of Cystinuria in a cohort of 12 patients. Clinical Genetics, 2012, 81, 47-55.	2.0	35
11	Mutational screening of VSX1, SPARC, SOD1, LOX, and TIMP3 in keratoconus. Molecular Vision, 2011, 17, 2482-94.	1.1	76
12	A new locus on 3p23-p25 for an autosomal-dominant limb-girdle muscular dystrophy, LGMD1H. European Journal of Human Genetics, 2010, 18, 636-641.	2.8	27
13	Large rearrangements detected by MLPA, point mutations, and survey of the frequency of mutations within the SLC3A1 and SLC7A9 genes in a cohort of 172 cystinuric Italian patients. Molecular Genetics and Metabolism, 2010, 99, 42-52.	1.1	34
14	Linkage Analysis in Keratoconus: Replication of Locus 5q21.2 and Identification of Other Suggestive Loci. , 2009, 50, 1081.		97
15	Molecular analysis in a family presenting with a mild form of late-onset autosomal dominant chronic progressive external ophthalmoplegia. Neuromuscular Disorders, 2009, 19, 423-426.	0.6	8
16	Genetic variant of C1GalT1 contributes to the susceptibility to IgA nephropathy. Journal of Nephrology, 2009, 22, 152-9.	2.0	30
17	An overview of SLC3A1 and SLC7A9 mutations in Greek cystinuria patients. Molecular Genetics and Metabolism, 2008, 95, 192-193.	1.1	9
18	Twenty-Four Novel Mutations Identified in a Cohort of 85 Patients by Direct Sequencing of the <i>SLC3A1</i> and <i>SLC7A9</i> Cystinuria Genes. Genetic Testing and Molecular Biomarkers, 2008, 12, 351-355.	1.7	16

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19	The Missing ApoE Allele. <i>Annals of Human Genetics</i> , 2007, 71, 496-500.	0.8	21
20	Human Gene Mutations. <i>Human Genetics</i> , 2007, 122, 207-215.	3.8	0
21	Genetic Heterogeneity in Italian Families with IgA Nephropathy: Suggestive Linkage for Two Novel IgA Nephropathy Loci. <i>American Journal of Human Genetics</i> , 2006, 79, 1130-1134.	6.2	111
22	Role of interferon- $\beta$ gene polymorphisms in susceptibility to IgA nephropathy: a family-based association study. <i>European Journal of Human Genetics</i> , 2006, 14, 488-496.	2.8	43
23	WT1 mutations in nephrotic syndrome revisited. High prevalence in young girls, associations and renal phenotypes. <i>Pediatric Nephrology</i> , 2006, 21, 1393-1398.	1.7	46
24	Posttransplant Recurrence of Proteinuria in a Case of Focal Segmental Glomerulosclerosis Associated with WT1 Mutation. <i>American Journal of Transplantation</i> , 2006, 6, 2208-2211.	4.7	19
25	Expression of VSX1 in Human Corneal Keratocytes during Differentiation into Myofibroblasts in Response to Wound Healing. , 2006, 47, 5243.		53
26	The IgA nephropathy Biobank. An important starting point for the genetic dissection of a complex trait. <i>BMC Nephrology</i> , 2005, 6, 14.	1.8	24
27	New insights into cystinuria: 40 new mutations, genotype-phenotype correlation, and digenic inheritance causing partial phenotype. <i>Journal of Medical Genetics</i> , 2005, 42, 58-68.	3.2	140
28	VSX1 Mutational Analysis in a Series of Italian Patients Affected by Keratoconus: Detection of a Novel Mutation. , 2005, 46, 39.		109
29	Indirect CFTR Mutation Identification by PCR/OLA Anomalous Electropherograms. <i>Genetic Testing and Molecular Biomarkers</i> , 2005, 9, 285-291.	1.7	1
30	Molecular Analysis of <i>NPHS2</i> and <i>ACTN4</i> Genes in a Series of 33 Italian Patients Affected by Adult-Onset Nonfamilial Focal Segmental Glomerulosclerosis. <i>Nephron Clinical Practice</i> , 2005, 99, c31-c36.	2.3	33
31	Comparison between SLC3A1 and SLC7A9 Cystinuria Patients and Carriers. <i>Journal of the American Society of Nephrology: JASN</i> , 2002, 13, 2547-2553.	6.1	234
32	MYO6, the Human Homologue of the Gene Responsible for Deafness in Snell's Waltzer Mice, Is Mutated in Autosomal Dominant Nonsyndromic Hearing Loss. <i>American Journal of Human Genetics</i> , 2001, 69, 635-640.	6.2	212
33	Cystinuria type I: Identification of eight new mutations in SLC3A1. <i>Kidney International</i> , 2001, 59, 1250-1256.	5.2	24
34	The putative forkhead transcription factor FOXL2 is mutated in blepharophimosis/ptosis/epicanthus inversus syndrome. <i>Nature Genetics</i> , 2001, 27, 159-166.	21.4	886
35	Functional analysis of mutations in SLC7A9, and genotype-phenotype correlation in non-Type I cystinuria. <i>Human Molecular Genetics</i> , 2001, 10, 305-316.	2.9	125
36	Detection of two novel large deletions in SLC3A1 by semi-quantitative fluorescent multiplex PCR. <i>Human Mutation</i> , 2000, 15, 373-379.	2.5	24

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37	Non-type I cystinuria caused by mutations in SLC7A9, encoding a subunit (bo,+AT) of rBAT. Nature Genetics, 1999, 23, 52-57.	21.4	280
38	Recombinant Families Locate the Gene for Non-Type I Cystinuria between Markers C13 and D19S587 on Chromosome 19q13.1. Genomics, 1999, 60, 362-365.	2.9	9
39	Polymorphism of motilin gene in patients with Crohn's disease. Digestive Diseases and Sciences, 1998, 43, 715-719.	2.3	7
40	Cellular Retinol Binding Protein 1 (RBP1): A frequent polymorphism, refined map position and exclusion as the Blepharophimosis Ptosis Epicanthus inversus Syndrome gene. Molecular and Cellular Probes, 1998, 12, 255-258.	2.1	2
41	Detection of dystrophin deletion carriers using FISH analysis. Clinical Genetics, 1997, 52, 17-22.	2.0	9
42	Genetic history of cystic fibrosis mutations in Italy. I. Regional distribution. Annals of Human Genetics, 1997, 61, 411-424.	0.8	26
43	A YAC Contig Spanning the Blepharophimosis-Ptosis- Epicanthus inversus Syndrome and Propionic Acidemia Loci. European Journal of Human Genetics, 1997, 5, 171-174.	2.8	7
44	Hereditary Hyperferritinemia-Cataract Syndrome Caused by a 29-Base Pair Deletion in the Iron Responsive Element of Ferritin L-Subunit Gene. Blood, 1997, 90, 2084-2088.	1.4	54
45	Genomic Structure and Organization of the Human rBAT Gene (SLC3A1). Genomics, 1996, 37, 249-252.	2.9	22
46	Molecular analysis of the cystinuria disease gene: identification of four new mutations, one large deletion, and one polymorphism. Human Genetics, 1996, 98, 447-451.	3.8	44
47	Phenotype characterization and prevalence of rBAT M467T mutation in Italian cystinuric patients. Journal of Inherited Metabolic Disease, 1996, 19, 243-245.	3.6	5
48	The molecular basis of cystinuria: the role of the rBAT gene. Amino Acids, 1996, 11, 225-246.	2.7	13
49	Genetic heterogeneity in cystinuria: the SLC3A1 gene is linked to type I but not to type III cystinuria.. Proceedings of the National Academy of Sciences of the United States of America, 1995, 92, 9667-9671.	7.1	98
50	Analysis of the complete coding region of the CFTR gene in a cohort of CF patients from North-Eastern Italy: identification of 90% of the mutations. Human Genetics, 1995, 95, 397-402.	3.8	56
51	Stability and functional effectiveness of phosphorothioate modified duplex DNA and synthetic $\hat{\text{a}}^{\text{c}}$ mini-genes $\hat{\text{a}}^{\text{TM}}$ . Nucleic Acids Research, 1995, 23, 4134-4142.	14.5	27
52	Molecular screening of genetic defects with RNA $\hat{\text{c}}$ SSCP analysis: the PKU and cystinuria model. Molecular and Cellular Probes, 1995, 9, 201-205.	2.1	3
53	Homozygosity for a novel splice site mutation (2790-2 A $\hat{\text{a}}$ 'G) preceding exon 15 of the CFTR gene in a cystic fibrosis patient of North-East Italian descent. Molecular and Cellular Probes, 1995, 9, 139-141.	2.1	1
54	The motilin gene: subregional localisation, tissue expression, DNA polymorphisms and exclusion as a candidate gene for the HLA-associated immotile cilia syndrome. Human Genetics, 1994, 94, 671-4.	3.8	15

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55	Development of RNA-SSCP protocols for the identification and screening of CFTR mutations: Identification of two new mutations. Human Mutation, 1994, 4, 136-140.	2.5	27