Luigi Bisceglia

List of Publications by Year in descending order

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		257450	1	182427	
55	3,367 citations	24		51	
papers	citations	h-index		g-index	
55	55	55		3618	
all docs	docs citations	times ranked		citing authors	

#	Article	IF	CITATIONS
1	The putative forkhead transcription factor FOXL2 is mutated in blepharophimosis/ptosis/epicanthus inversus syndrome. Nature Genetics, 2001, 27, 159-166.	21.4	886
2	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
3	Comparison between SLC3A1 and SLC7A9 Cystinuria Patients and Carriers. Journal of the American Society of Nephrology: JASN, 2002, 13, 2547-2553.	6.1	234
4	MYO6, the Human Homologue of the Gene Responsible for Deafness in Snell's Waltzer Mice, Is Mutated in Autosomal Dominant Nonsyndromic Hearing Loss. American Journal of Human Genetics, 2001, 69, 635-640.	6.2	212
5	New insights into cystinuria: 40 new mutations, genotype-phenotype correlation, and digenic inheritance causing partial phenotype. Journal of Medical Genetics, 2005, 42, 58-68.	3.2	140
6	Functional analysis of mutations in SLC7A9, and genotype–phenotype correlation in non-Type I cystinuria. Human Molecular Genetics, 2001, 10, 305-316.	2.9	125
7	Genetic Heterogeneity in Italian Families with IgA Nephropathy: Suggestive Linkage for Two Novel IgA Nephropathy Loci. American Journal of Human Genetics, 2006, 79, 1130-1134.	6.2	111
8	VSX1Mutational Analysis in a Series of Italian Patients Affected by Keratoconus: Detection of a Novel Mutation., 2005, 46, 39.		109
9	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
10	Linkage Analysis in Keratoconus: Replication of Locus 5q21.2 and Identification of Other Suggestive Loci., 2009, 50, 1081.		97
11	Mutational screening of VSX1, SPARC, SOD1, LOX, and TIMP3 in keratoconus. Molecular Vision, 2011, 17, 2482-94.	1.1	76
12	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
13	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
14	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
15	Expression of VSX1 in Human Corneal Keratocytes during Differentiation into Myofibroblasts in Response to Wound Healing., 2006, 47, 5243.		53
16	WT1 mutations in nephrotic syndrome revisited. High prevalence in young girls, associations and renal phenotypes. Pediatric Nephrology, 2006, 21, 1393-1398.	1.7	46
17	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
18	Role of interferon- \hat{l}^3 gene polymorphisms in susceptibility to IgA nephropathy: a family-based association study. European Journal of Human Genetics, 2006, 14, 488-496.	2.8	43

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19	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
20	Clinical, biochemical and molecular characterization of Cystinuria in a cohort of 12 patients. Clinical Genetics, 2012, 81, 47-55.	2.0	35
21	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
22	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. Nephron Clinical Practice, 2005, 99, c31-c36.	2.3	33
23	Genetic variant of C1GalT1 contributes to the susceptibility to IgA nephropathy. Journal of Nephrology, 2009, 22, 152-9.	2.0	30
24	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
25	Stability and functional effectiveness of phosphorothioate modified duplex DNA and synthetic â€~mini-genes'. Nucleic Acids Research, 1995, 23, 4134-4142.	14.5	27
26	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
27	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
28	Genetic history of cystic fibrosis mutations in Italy. I. Regional distribution. Annals of Human Genetics, 1997, 61, 411-424.	0.8	26
29	Mitochondrial DNA copy number in affected and unaffected LHON mutation carriers. BMC Research Notes, 2018, 11, 911.	1.4	25
30	Detection of two novel large deletions in SLC3A1 by semi-quantitative fluorescent multiplex PCR. Human Mutation, 2000, 15, 373-379.	2.5	24
31	Cystinuria type I: Identification of eight new mutations in SLC3A1. Kidney International, 2001, 59, 1250-1256.	5.2	24
32	The IgA nephropathy Biobank. An important starting point for the genetic dissection of a complex trait. BMC Nephrology, 2005, 6, 14.	1.8	24
33	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
34	Genomic Structure and Organization of the HumanrBATGene (SLC3A1). Genomics, 1996, 37, 249-252.	2.9	22
35	The Missing ApoE Allele. Annals of Human Genetics, 2007, 71, 496-500.	0.8	21
36	Posttransplant Recurrence of Proteinuria in a Case of Focal Segmental Glomerulosclerosis Associated with WT1 Mutation. American Journal of Transplantation, 2006, 6, 2208-2211.	4.7	19

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37	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
38	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
39	Identification and Clinical Characterization of Adult Patients with Multigenerational Diabetes Mellitus. PLoS ONE, 2015, 10, e0135855.	2.5	14
40	The molecular basis of cystinuria: the role of the rBAT gene. Amino Acids, 1996, 11, 225-246.	2.7	13
41	Clinical utility gene card for: Cystinuria. European Journal of Human Genetics, 2012, 20, 3-3.	2.8	12
42	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
43	Detection of dystrophin deletion carriers using FISH analysis. Clinical Genetics, 1997, 52, 17-22.	2.0	9
44	An overview of SLC3A1 and SLC7A9 mutations in Greek cystinuria patients. Molecular Genetics and Metabolism, 2008, 95, 192-193.	1.1	9
45	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
46	Polymorphism of motilin gene in patients with Crohn's disease. Digestive Diseases and Sciences, 1998, 43, 715-719.	2.3	7
47	Leber's hereditary optic neuropathy, intellectual disability and epilepsy presenting with variable penetrance associated to the m.3460G >A mutation and a heteroplasmic expansion of the microsatellite in MTRNR1 gene – case report. BMC Medical Genetics, 2018, 19, 129.	2.1	7
48	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
49	Leber's hereditary optic neuropathy (LHON) in an Apulian cohort of subjects. Acta Myologica, 2017, 36, 163-177.	1.5	7
50	Phenotype characterization and prevalence of rBAT M467T mutation in Italian cystinuric patients. Journal of Inherited Metabolic Disease, 1996, 19, 243-245.	3.6	5
51	Molecular screening of genetic defects with RNA–SSCP analysis: the PKU and cystinuria model. Molecular and Cellular Probes, 1995, 9, 201-205.	2.1	3
52	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
53	Homozygosity for a novel splice site mutation (2790-2 Aâ†'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	IndirectCFTRMutation Identification by PCR/OLA Anomalous Electropherograms. Genetic Testing and Molecular Biomarkers, 2005, 9, 285-291.	1.7	1

#	Article	IF	CITATIONS
55	Human Gene Mutations. Human Genetics, 2007, 122, 207-215.	3.8	0