

Dominique Bonneau

List of Publications by Year in descending order

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Version: 2024-02-01

186
papers

13,372
citations

31976

53
h-index

25787

108
g-index

196
all docs

196
docs citations

196
times ranked

17932
citing authors

#	ARTICLE	IF	CITATIONS
1	Glutamate-Induced Deregulation of Krebs Cycle in Mitochondrial Encephalopathy Lactic Acidosis Syndrome Stroke-Like Episodes (MELAS) Syndrome Is Alleviated by Ketone Body Exposure. <i>Biomedicines</i> , 2022, 10, 1665.	3.2	4
2	Next-Generation Sequencing Identifies Novel PMPCA Variants in Patients with Late-Onset Dominant Optic Atrophy. <i>Genes</i> , 2022, 13, 1202.	2.4	0
3	Prenatal diagnosis of Desbuquois dysplasia type 1 by whole exome sequencing before the occurrence of specific ultrasound signs. <i>Journal of Maternal-Fetal and Neonatal Medicine</i> , 2021, 34, 2217-2220.	1.5	5
4	A plasma metabolomic signature of Leber hereditary optic neuropathy showing taurine and nicotinamide deficiencies. <i>Human Molecular Genetics</i> , 2021, 30, 21-29.	2.9	14
5	Dominant mutations in MIEF1 affect mitochondrial dynamics and cause a singular late onset optic neuropathy. <i>Molecular Neurodegeneration</i> , 2021, 16, 12.	10.8	13
6	Use of Next-Generation Sequencing for the Molecular Diagnosis of 1,102 Patients With a Autosomal Optic Neuropathy. <i>Frontiers in Neurology</i> , 2021, 12, 602979.	2.4	12
7	Metabolomic Sexual Dimorphism of the Mouse Brain is Predominantly Abolished by Gonadectomy with a Higher Impact on Females. <i>Journal of Proteome Research</i> , 2021, 20, 2772-2779.	3.7	1
8	Are Your Mitochondria Ready for a Space Odyssey?. <i>Trends in Endocrinology and Metabolism</i> , 2021, 32, 193-195.	7.1	1
9	Dominant <i>ACO2</i> mutations are a frequent cause of isolated optic atrophy. <i>Brain Communications</i> , 2021, 3, fcab063.	3.3	16
10	Improved detection of mitochondrial DNA instability in mitochondrial genome maintenance disorders. <i>Genetics in Medicine</i> , 2021, 23, 1769-1778.	2.4	4
11	Missense variants in <i>DPYSL5</i> cause a neurodevelopmental disorder with corpus callosum agenesis and cerebellar abnormalities. <i>American Journal of Human Genetics</i> , 2021, 108, 951-961.	6.2	26
12	Bi-allelic variants in <i>IPO8</i> cause a connective tissue disorder associated with cardiovascular defects, skeletal abnormalities, and immune dysregulation. <i>American Journal of Human Genetics</i> , 2021, 108, 1126-1137.	6.2	14
13	Heterozygous <i>HMGB1</i> loss-of-function variants are associated with developmental delay and microcephaly. <i>Clinical Genetics</i> , 2021, 100, 386-395.	2.0	3
14	<i>ZNF668</i> deficiency causes a recognizable disorder of DNA damage repair. <i>Human Genetics</i> , 2021, 140, 1395-1401.	3.8	1
15	Psychiatric Symptoms of Children and Adolescents With Mitochondrial Disorders: A Descriptive Case Series. <i>Frontiers in Psychiatry</i> , 2021, 12, 685532.	2.6	1
16	Mobile App for Parental Empowerment for Caregivers of Children With Autism Spectrum Disorders: Prospective Open Trial. <i>JMIR Mental Health</i> , 2021, 8, e27803.	3.3	7
17	Congenital hypothyroidism and hearing loss without inner ear malformation: Think <i>TPO</i> . <i>Clinical Genetics</i> , 2021, 99, 604-606.	2.0	1
18	Metabolomics reveals highly regional specificity of cerebral sexual dimorphism in mice. <i>Progress in Neurobiology</i> , 2020, 184, 101698.	5.7	11

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19	De novo and inherited variants in ZNF292 underlie a neurodevelopmental disorder with features of autism spectrum disorder. <i>Genetics in Medicine</i> , 2020, 22, 538-546.	2.4	24
20	Oxidative stress contributes differentially to the pathophysiology of Charcot-Marie-Tooth disease type 2K. <i>Experimental Neurology</i> , 2020, 323, 113069.	4.1	22
21	Specific cognitive theory of mind and behavioral dysfunctions in early manifest Huntington disease: a case report. <i>Neurocase</i> , 2020, 26, 36-41.	0.6	1
22	Phenotypic spectrum of <i>TGFB3</i> disease-causing variants in a Dutch-French cohort and first report of a homozygous patient. <i>Clinical Genetics</i> , 2020, 97, 723-730.	2.0	15
23	Germline and Mosaic Variants in PRKACA and PRKACB Cause a Multiple Congenital Malformation Syndrome. <i>American Journal of Human Genetics</i> , 2020, 107, 977-988.	6.2	33
24	Dysfunctional T Cell Mitochondria Lead to Premature Aging. <i>Trends in Molecular Medicine</i> , 2020, 26, 799-800.	6.7	5
25	Clinical and molecular description of 19 patients with GATAD2B-Associated Neurodevelopmental Disorder (GAND). <i>European Journal of Medical Genetics</i> , 2020, 63, 104004.	1.3	7
26	Neuropsychological and Psychiatric Features of Children and Adolescents Affected With Mitochondrial Diseases: A Systematic Review. <i>Frontiers in Psychiatry</i> , 2020, 11, 747.	2.6	12
27	De novo missense variants in LMBRD2 are associated with developmental and motor delays, brain structure abnormalities and dysmorphic features. <i>Journal of Medical Genetics</i> , 2020, 58, jmedgenet-2020-107137.	3.2	3
28	Hypopituitarism in Patients with Blepharophimosis and <i>FOXL2</i> Mutations. <i>Hormone Research in Paediatrics</i> , 2020, 93, 30-39.	1.8	8
29	Metabolomics hallmarks OPA1 variants correlating with their in vitro phenotype and predicting clinical severity. <i>Human Molecular Genetics</i> , 2020, 29, 1319-1329.	2.9	17
30	A Data Mining Metabolomics Exploration of Glaucoma. <i>Metabolites</i> , 2020, 10, 49.	2.9	25
31	A new mutational hotspot in the SKI gene in the context of MFS/TAA molecular diagnosis. <i>Human Genetics</i> , 2020, 139, 461-472.	3.8	8
32	Mutations in aARS genes revealed by targeted next-generation sequencing in patients with mitochondrial diseases. <i>Molecular Biology Reports</i> , 2020, 47, 3779-3787.	2.3	5
33	Mutations in the m-AAA proteases AFG3L2 and SPG7 are causing isolated dominant optic atrophy. <i>Neurology: Genetics</i> , 2020, 6, e428.	1.9	31
34	AMPA receptor GluA2 subunit defects are a cause of neurodevelopmental disorders. <i>Nature Communications</i> , 2019, 10, 3094.	12.8	150
35	Confirmation that variants in <i>TTI2</i> are responsible for autosomal recessive intellectual disability. <i>Clinical Genetics</i> , 2019, 96, 354-358.	2.0	14
36	CTCF variants in 39 individuals with a variable neurodevelopmental disorder broaden the mutational and clinical spectrum. <i>Genetics in Medicine</i> , 2019, 21, 2723-2733.	2.4	48

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37	Developmental trajectories of neuroanatomical alterations associated with the 16p11.2 Copy Number Variations. <i>NeuroImage</i> , 2019, 203, 116155.	4.2	9
38	Metabolomic Profiling of Aqueous Humor in Glaucoma Points to Taurine and Spermine Deficiency: Findings from the Eye-D Study. <i>Journal of Proteome Research</i> , 2019, 18, 1307-1315.	3.7	49
39	Nicotinamide Deficiency in Primary Open-Angle Glaucoma. , 2019, 60, 2509.		61
40	Lipidomics Reveals Triacylglycerol Accumulation Due to Impaired Fatty Acid Flux in <i>OPA1</i> -Disrupted Fibroblasts. <i>Journal of Proteome Research</i> , 2019, 18, 2779-2790.	3.7	9
41	Warburg-like effect is a hallmark of complex I assembly defects. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2019, 1865, 2475-2489.	3.8	13
42	Where are the missing gene defects in inherited retinal disorders? Intronic and synonymous variants contribute at least to 4% of <i>CACNA1F</i> -mediated inherited retinal disorders. <i>Human Mutation</i> , 2019, 40, 765-787.	2.5	24
43	A snapshot of some pLI score pitfalls. <i>Human Mutation</i> , 2019, 40, 839-841.	2.5	29
44	A New Case of <i>PCSK1</i> Pathogenic Variant With Congenital Proprotein Convertase 1/3 Deficiency and Literature Review. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2019, 104, 985-993.	3.6	39
45	Missense Variants in the Histone Acetyltransferase Complex Component Gene <i>TRRAP</i> Cause Autism and Syndromic Intellectual Disability. <i>American Journal of Human Genetics</i> , 2019, 104, 530-541.	6.2	30
46	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
47	Next generation sequencing in family with MNGIE syndrome associated to optic atrophy: Novel homozygous <i>POLG</i> mutation in the C-terminal sub-domain leading to mtDNA depletion. <i>Clinica Chimica Acta</i> , 2019, 488, 104-110.	1.1	12
48	Phenotypic spectrum associated with <i>SPECC1L</i> pathogenic variants: new families and critical review of the nosology of Teebi, Opitz GBBB, and Baraitser-Winter syndromes. <i>European Journal of Medical Genetics</i> , 2019, 62, 103588.	1.3	24
49	eKLIPse: a sensitive tool for the detection and quantification of mitochondrial DNA deletions from next-generation sequencing data. <i>Genetics in Medicine</i> , 2019, 21, 1407-1416.	2.4	38
50	Expanding the phenotypic spectrum associated with <i>OPHN1</i> mutations: Report of 17 individuals with intellectual disability but no cerebellar hypoplasia. <i>European Journal of Medical Genetics</i> , 2018, 61, 442-450.	1.3	8
51	Quantifying the Effects of 16p11.2 Copy Number Variants on Brain Structure: A Multisite Genetic-First Study. <i>Biological Psychiatry</i> , 2018, 84, 253-264.	1.3	56
52	The accumulation of assembly intermediates of the mitochondrial complex I matrix arm is reduced by limiting glucose uptake in a neuronal-like model of MELAS syndrome. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 1596-1608.	3.8	15
53	Reply: The expanding neurological phenotype of <i>DNM1L</i> -related disorders. <i>Brain</i> , 2018, 141, e29-e29.	7.6	5
54	Neurologic Phenotypes Associated With Mutations in <i>RTN4IP1</i> (<i>OPA10</i>) in Children and Young Adults. <i>JAMA Neurology</i> , 2018, 75, 105.	9.0	26

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55	Spastic paraplegia due to SPAST mutations is modified by the underlying mutation and sex. <i>Brain</i> , 2018, 141, 3331-3342.	7.6	72
56	Bioinformatics Tools and Databases to Assess the Pathogenicity of Mitochondrial DNA Variants in the Field of Next Generation Sequencing. <i>Frontiers in Genetics</i> , 2018, 9, 632.	2.3	48
57	A Plasma Metabolomic Signature Involving Purine Metabolism in Human Optic Atrophy 1 (<i>OPA1</i>)-Related Disorders. , 2018, 59, 185.		21
58	A Metabolomics Profiling of Glaucoma Points to Mitochondrial Dysfunction, Senescence, and Polyamines Deficiency. , 2018, 59, 4355.		51
59	A Plasma Metabolomic Signature of the Exfoliation Syndrome Involves Amino Acids, Acylcarnitines, and Polyamines. , 2018, 59, 1025.		18
60	The Metabolomic Bioenergetic Signature of Opa1-Disrupted Mouse Embryonic Fibroblasts Highlights Aspartate Deficiency. <i>Scientific Reports</i> , 2018, 8, 11528.	3.3	20
61	De Novo Disruption of the Proteasome Regulatory Subunit PSMD12 Causes a Syndromic Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2017, 100, 352-363.	6.2	86
62	Haploinsufficiency of the E3 ubiquitin-protein ligase gene TRIP12 causes intellectual disability with or without autism spectrum disorders, speech delay, and dysmorphic features. <i>Human Genetics</i> , 2017, 136, 377-386.	3.8	36
63	Gain-of-Function Mutation in Filamin A Potentiates Platelet Integrin α IIb β 3 Activation. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2017, 37, 1087-1097.	2.4	28
64	CLUH couples mitochondrial distribution to the energetic and metabolic status. <i>Journal of Cell Science</i> , 2017, 130, 1940-1951.	2.0	38
65	A randomized, double-blind, placebo-controlled trial evaluating cysteamine in Huntington's disease. <i>Movement Disorders</i> , 2017, 32, 932-936.	3.9	31
66	Identification of genetic variants associated with Huntington's disease progression: a genome-wide association study. <i>Lancet Neurology</i> , The, 2017, 16, 701-711.	10.2	248
67	Autophagy controls the pathogenicity of <i>OPA1</i> mutations in dominant optic atrophy. <i>Journal of Cellular and Molecular Medicine</i> , 2017, 21, 2284-2297.	3.6	30
68	A Nontargeted UHPLC-HRMS Metabolomics Pipeline for Metabolite Identification: Application to Cardiac Remote Ischemic Preconditioning. <i>Analytical Chemistry</i> , 2017, 89, 2138-2146.	6.5	43
69	De Novo Missense Mutations in DHX30 Impair Global Translation and Cause a Neurodevelopmental Disorder. <i>American Journal of Human Genetics</i> , 2017, 101, 716-724.	6.2	66
70	De Novo Mutations in Protein Kinase Genes CAMK2A and CAMK2B Cause Intellectual Disability. <i>American Journal of Human Genetics</i> , 2017, 101, 768-788.	6.2	136
71	Mutations in DNM1L, as in OPA1, result in dominant optic atrophy despite opposite effects on mitochondrial fusion and fission. <i>Brain</i> , 2017, 140, 2586-2596.	7.6	100
72	A recessive ataxia diagnosis algorithm for the next generation sequencing era. <i>Annals of Neurology</i> , 2017, 82, 892-899.	5.3	27

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73	Dissociation between decision-making under risk and decision-making under ambiguity in premanifest and manifest Huntington's disease. <i>Neuropsychologia</i> , 2017, 103, 87-95.	1.6	11
74	The addition of ketone bodies alleviates mitochondrial dysfunction by restoring complex I assembly in a MELAS cellular model. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2017, 1863, 284-291.	3.8	41
75	Novel <i>NDUFS4</i> gene mutation in an atypical late-onset mitochondrial form of multifocal dystonia. <i>Neurology: Genetics</i> , 2017, 3, e205.	1.9	7
76	Targeted Metabolomics Reveals Early Dominant Optic Atrophy Signature in Optic Nerves of <i>OPA1</i> ^{ΔTTAG/+} Mice. , 2017, 58, 812.		22
77	Rationale and protocol for using a smartphone application to study autism spectrum disorders: SMARTAUTISM. <i>BMJ Open</i> , 2016, 6, e012135.	1.9	5
78	Biallelic Variants in UBA5 Reveal that Disruption of the UFM1 Cascade Can Result in Early-Onset Encephalopathy. <i>American Journal of Human Genetics</i> , 2016, 99, 695-703.	6.2	87
79	The metabolomic signature of Leber's hereditary optic neuropathy reveals endoplasmic reticulum stress. <i>Brain</i> , 2016, 139, 2864-2876.	7.6	45
80	Increased mitochondrial fusion in a autosomal recessive CMT2A family with mitochondrial GTPase mitofusin 2 mutations. <i>Journal of the Peripheral Nervous System</i> , 2016, 21, 365-369.	3.1	10
81	De Novo Truncating Mutations in the Kinetochores-Microtubules Attachment Gene <i>CHAMP1</i> Cause Syndromic Intellectual Disability. <i>Human Mutation</i> , 2016, 37, 354-358.	2.5	40
82	Run of homozygosity analysis reveals a novel nonsense variant of the <i>CNGB1</i> gene involved in retinitis pigmentosa 45. <i>Ophthalmic Genetics</i> , 2016, 37, 357-359.	1.2	7
83	OPA1-related disorders: Diversity of clinical expression, modes of inheritance and pathophysiology. <i>Neurobiology of Disease</i> , 2016, 90, 20-26.	4.4	45
84	Theory of mind and empathy in preclinical and clinical Huntington's disease. <i>Social Cognitive and Affective Neuroscience</i> , 2016, 11, 89-99.	3.0	27
85	New practical definitions for the diagnosis of autosomal recessive spastic ataxia of Charlevoix-Saguenay. <i>Annals of Neurology</i> , 2015, 78, 871-886.	5.3	62
86	Improved Locus-Specific Database for <i>OPA1</i> Mutations Allows Inclusion of Advanced Clinical Data. <i>Human Mutation</i> , 2015, 36, 20-25.	2.5	39
87	A phase II, open-label evaluation of cysteamine tolerability in patients with Huntington's disease. <i>Movement Disorders</i> , 2015, 30, 288-289.	3.9	16
88	Assembly defects induce oxidative stress in inherited mitochondrial complex I deficiency. <i>International Journal of Biochemistry and Cell Biology</i> , 2015, 65, 91-103.	2.8	29
89	Adult-onset genetic leukoencephalopathies: A MRI pattern-based approach in a comprehensive study of 154 patients. <i>Brain</i> , 2015, 138, 284-292.	7.6	58
90	Recessive Mutations in <i>RTN4IP1</i> Cause Isolated and Syndromic Optic Neuropathies. <i>American Journal of Human Genetics</i> , 2015, 97, 754-760.	6.2	54

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91	Dermatologic Features of Smithâ€“Magenis Syndrome. <i>Pediatric Dermatology</i> , 2015, 32, 337-341.	0.9	10
92	Mutations in CNTNAP1 and ADCY6 are responsible for severe arthrogyriposis multiplex congenita with axogial defects. <i>Human Molecular Genetics</i> , 2014, 23, 2279-2289.	2.9	98
93	Meta-analysis of SHANK Mutations in Autism Spectrum Disorders: A Gradient of Severity in Cognitive Impairments. <i>PLoS Genetics</i> , 2014, 10, e1004580.	3.5	501
94	Early-onset Behr syndrome due to compound heterozygous mutations in OPA1. <i>Brain</i> , 2014, 137, e301-e301.	7.6	62
95	Loss-of-Function Mutations in WDR73 Are Responsible for Microcephaly and Steroid-Resistant Nephrotic Syndrome: Galloway-Mowat Syndrome. <i>American Journal of Human Genetics</i> , 2014, 95, 637-648.	6.2	108
96	Enrichment of LOVD-USHbases with 152<i>USH2A</i> Genotypes Defines an Extensive Mutational Spectrum and Highlights Missense Hotspots. <i>Human Mutation</i> , 2014, 35, 1179-1186.	2.5	55
97	Extensive Mongolian spots in 4p16.3 deletion (Wolfâ€“Hirschhorn syndrome). <i>Clinical Dysmorphology</i> , 2014, 23, 109-110.	0.3	2
98	SLC24A5 Mutations Are Associated with Non-Syndromic Oculocutaneous Albinism. <i>Journal of Investigative Dermatology</i> , 2014, 134, 568-571.	0.7	36
99	Efficient strategy for the molecular diagnosis of intellectual disability using targeted high-throughput sequencing. <i>Journal of Medical Genetics</i> , 2014, 51, 724-736.	3.2	229
100	Evaluation of the colorectal cancer risk conferred by rare<i>UNC5C</i> alleles. <i>World Journal of Gastroenterology</i> , 2014, 20, 204.	3.3	11
101	Mutations in TUBG1, DYNC1H1, KIF5C and KIF2A cause malformations of cortical development and microcephaly. <i>Nature Genetics</i> , 2013, 45, 639-647.	21.4	399
102	Resveratrol Induces a Mitochondrial Complex I-dependent Increase in NADH Oxidation Responsible for Sirtuin Activation in Liver Cells. <i>Journal of Biological Chemistry</i> , 2013, 288, 36662-36675.	3.4	110
103	Relative Frequencies of Inherited Retinal Dystrophies and Optic Neuropathies in Southern France: Assessment of 21-year Data Management. <i>Ophthalmic Epidemiology</i> , 2013, 20, 13-25.	1.7	44
104	High cumulative risks of cancer in patients with<i>PTEN</i> hamartoma tumour syndrome. <i>Journal of Medical Genetics</i> , 2013, 50, 255-263.	3.2	290
105	Sensorineural hearing loss in OPA1-linked disorders. <i>Brain</i> , 2013, 136, e236-e236.	7.6	36
106	Intellectual disability associated with retinal dystrophy in the Xp11.3 deletion syndrome: ZNF674 on trial. Guilty or innocent?. <i>European Journal of Human Genetics</i> , 2012, 20, 352-356.	2.8	5
107	Genetic and Functional Analyses of SHANK2 Mutations Suggest a Multiple Hit Model of Autism Spectrum Disorders. <i>PLoS Genetics</i> , 2012, 8, e1002521.	3.5	358
108	Cobblestone lissencephaly: neuropathological subtypes and correlations with genes of dystroglycanopathies. <i>Brain</i> , 2012, 135, 469-482.	7.6	151

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109	Metabolically induced heteroplasmy shifting and l-arginine treatment reduce the energetic defect in a neuronal-like model of MELAS. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2012, 1822, 1019-1029.	3.8	38
110	Standardized mitochondrial analysis gives new insights into mitochondrial dynamics and OPA1 function. <i>International Journal of Biochemistry and Cell Biology</i> , 2012, 44, 980-988.	2.8	34
111	Dominant optic atrophy. <i>Orphanet Journal of Rare Diseases</i> , 2012, 7, 46.	2.7	213
112	Reduced elastogenesis: a clue to the arteriosclerosis and emphysematous changes in Schimke immuno-osseous dysplasia?. <i>Orphanet Journal of Rare Diseases</i> , 2012, 7, 70.	2.7	26
113	Prenatal diagnosis of CHARGE syndrome by identification of a novel <i>CHD7</i> mutation in a previously unaffected family. <i>Prenatal Diagnosis</i> , 2012, 32, 692-694.	2.3	15
114	Whole-Exome Sequencing Identifies Mutations in GPR179 Leading to Autosomal-Recessive Complete Congenital Stationary Night Blindness. <i>American Journal of Human Genetics</i> , 2012, 90, 321-330.	6.2	121
115	Spectrum of mutations in the renin-angiotensin system genes in autosomal recessive renal tubular dysgenesis. <i>Human Mutation</i> , 2012, 33, 316-326.	2.5	86
116	A <i>de novo</i> germline <i>MLH1</i> mutation in a Lynch syndrome patient with discordant immunohistochemical and molecular biology test results. <i>World Journal of Gastroenterology</i> , 2012, 18, 5635.	3.3	5
117	Mirror extreme BMI phenotypes associated with gene dosage at the chromosome 16p11.2 locus. <i>Nature</i> , 2011, 478, 97-102.	27.8	394
118	Molecular diagnosis reveals genetic heterogeneity for the overlapping MKKS and BBS phenotypes. <i>European Journal of Medical Genetics</i> , 2011, 54, 157-160.	1.3	39
119	Thrombocytopenia-absent radius (TAR) syndrome: A clinical genetic series of 14 further cases. Impact of the associated 1q21.1 deletion on the genetic counselling. <i>European Journal of Medical Genetics</i> , 2011, 54, e471-e477.	1.3	29
120	The neural substrates of script knowledge deficits as revealed by a PET study in Huntington's disease. <i>Neuropsychologia</i> , 2011, 49, 2673-2684.	1.6	9
121	Idebenone increases mitochondrial complex I activity in fibroblasts from LHON patients while producing contradictory effects on respiration. <i>BMC Research Notes</i> , 2011, 4, 557.	1.4	50
122	Hereditary spastic paraplegia-like disorder due to a mitochondrial ATP6 gene point mutation. <i>Mitochondrion</i> , 2011, 11, 70-75.	3.4	74
123	Mutation screening of the EYA1, SIX1, and SIX5 genes in a large cohort of patients harboring branchio-oto-renal syndrome calls into question the pathogenic role of SIX5 mutations. <i>Human Mutation</i> , 2011, 32, 183-190.	2.5	93
124	Bioenergetic defect associated with mK _{ATP} channel opening in a mouse model carrying a mitofusin 2 mutation. <i>FASEB Journal</i> , 2011, 25, 1618-1627.	0.5	36
125	Genetically determined optic neuropathies. <i>Current Opinion in Neurology</i> , 2010, 23, 24-28.	3.6	31
126	Adenine nucleotide translocase is involved in a mitochondrial coupling defect in MFN2-related Charcot-Marie-Tooth type 2A disease. <i>Neurogenetics</i> , 2010, 11, 127-133.	1.4	34

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127	Schimke immunoosseous dysplasia: defining skeletal features. <i>European Journal of Pediatrics</i> , 2010, 169, 801-811.	2.7	22
128	Search for the best indicators for the presence of a VPS13B gene mutation and confirmation of diagnostic criteria in a series of 34 patients genotyped for suspected Cohen syndrome. <i>Journal of Medical Genetics</i> , 2010, 47, 549-553.	3.2	31
129	Ethambutol-induced optic neuropathy linked to OPA1 mutation and mitochondrial toxicity. <i>Mitochondrion</i> , 2010, 10, 115-124.	3.4	49
130	Identification of gene copy number variations in patients with mental retardation using array-CGH: Novel syndromes in a large French series. <i>European Journal of Medical Genetics</i> , 2010, 53, 66-75.	1.3	29
131	Molecular screening of 980 cases of suspected hereditary optic neuropathy with a report on 77 novel OPA1 mutations. <i>Human Mutation</i> , 2009, 30, E692-E705.	2.5	140
132	Mitochondrial complex I deficiency in GDAP1-related autosomal dominant Charcot-Marie-Tooth disease (CMT2K). <i>Neurogenetics</i> , 2009, 10, 145-150.	1.4	72
133	A large-scale mutation search reveals genetic heterogeneity in 3M syndrome. <i>European Journal of Human Genetics</i> , 2009, 17, 395-400.	2.8	48
134	OPA1-associated disorders: Phenotypes and pathophysiology. <i>International Journal of Biochemistry and Cell Biology</i> , 2009, 41, 1855-1865.	2.8	122
135	Acute and late-onset optic atrophy due to a novel OPA1 mutation leading to a mitochondrial coupling defect. <i>Molecular Vision</i> , 2009, 15, 598-608.	1.1	37
136	Reversible optic neuropathy with OPA1 exon 5b mutation. <i>Annals of Neurology</i> , 2008, 63, 667-671.	5.3	36
137	Hereditary optic neuropathies share a common mitochondrial coupling defect. <i>Annals of Neurology</i> , 2008, 63, 794-798.	5.3	112
138	Molecular cytogenetic characterization of terminal 14q32 deletions in two children with an abnormal phenotype and corpus callosum hypoplasia. <i>European Journal of Human Genetics</i> , 2008, 16, 680-687.	2.8	23
139	OPA1 mutations induce mitochondrial DNA instability and optic atrophy 'plus' phenotypes. <i>Brain</i> , 2008, 131, 338-351.	7.6	454
140	Heterozygosity for a Single Mutation in the ABCC6 Gene May Closely Mimic PXE. <i>Archives of Dermatology</i> , 2008, 144, 301-6.	1.4	36
141	Identification of a Novel BBS Gene (BBS12) Highlights the Major Role of a Vertebrate-Specific Branch of Chaperonin-Related Proteins in Bardet-Biedl Syndrome. <i>American Journal of Human Genetics</i> , 2007, 80, 1-11.	6.2	219
142	Mitochondrial coupling defect in Charcot-Marie-Tooth type 2A disease. <i>Annals of Neurology</i> , 2007, 61, 315-323.	5.3	127
143	Schimke immunoosseous dysplasia: suggestions of genetic diversity. <i>Human Mutation</i> , 2007, 28, 273-283.	2.5	49
144	Molecular and in silico analyses of the full-length isoform of usherin identify new pathogenic alleles in Usher type II patients. <i>Human Mutation</i> , 2007, 28, 781-789.	2.5	98

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146	Effects of OPA1 mutations on mitochondrial morphology and apoptosis: Relevance to ADOA pathogenesis. <i>Journal of Cellular Physiology</i> , 2007, 211, 423-430.	4.1	128
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152	Pitfalls of homozygosity mapping: an extended consanguineous Bardet-Biedl syndrome family with two mutant genes (BBS2, BBS10), three mutations, but no triallelism. <i>European Journal of Human Genetics</i> , 2006, 14, 1195-1203.	2.8	56
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162	Mutations of ARX are associated with striking pleiotropy and consistent genotype-phenotype correlation. <i>Human Mutation</i> , 2004, 23, 147-159.	2.5	293

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