Borut Peterlin

List of Publications by Year in descending order

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		117625	114465
219	5,498	34	63
papers	citations	h-index	g-index
222	222	222	8815
all docs	docs citations	times ranked	citing authors

#	Article	lF	CITATIONS
1	Maternal LINE-1 DNA Methylation in Early Spontaneous Preterm Birth. Biological Research for Nursing, 2022, 24, 85-93.	1.9	2
2	A case report of a novel GNB1 pathogenic variant and the response to deep brain stimulation. Acta Neurologica Belgica, 2022, , 1.	1.1	3
3	Syndromic male subfertility: A network view of genome–phenome associations. Andrology, 2022, 10, 720-732.	3.5	5
4	Axonal Polyneuropathy in 2 Brothers With a Homozygous Missense Variant in the First Catalytic Domain of <i>PCYT2</i> . Neurology: Genetics, 2022, 8, e658.	1.9	2
5	Editorial: The Importance of Genetic Literacy and Education in Medicine. Frontiers in Genetics, 2022, 13, 910530.	2.3	0
6	Biallelic <i>ATOH1</i> Gene Variant in Siblings With Pontocerebellar Hypoplasia, Developmental Delay, and Hearing Loss. Neurology: Genetics, 2022, 8, e677.	1.9	2
7	Family history tools for primary care: A systematic review. European Journal of General Practice, 2022, 28, 75-86.	2.0	7
8	Clinical exome sequencing in Serbian patients with movement disorders: Single centre experience. Genetika, 2022, 54, 395-409.	0.4	1
9	The Endometrial Transcriptome of Metabolic and Inflammatory Pathways During the Window of Implantation Is Deranged in Infertile Obese Polycystic Ovarian Syndrome Women. Metabolic Syndrome and Related Disorders, 2022, 20, 384-394.	1.3	4
10	De novo mutations in idiopathic male infertility—A pilot study. Andrology, 2021, 9, 212-220.	3.5	19
11	Improving diagnostics of rare genetic diseases with NGS approaches. Journal of Community Genetics, 2021, 12, 247-256.	1.2	25
12	Stationary and Progressive Phenotypes Caused by the p.G90D Mutation in Rhodopsin Gene. International Journal of Molecular Sciences, 2021, 22, 2133.	4.1	6
13	Cone Dystrophy Associated with a Novel Variant in the Terminal Codon of the RPGR-ORF15. Genes, 2021, 12, 499.	2.4	8
14	How to design a national genomic project—a systematic review of active projects. Human Genomics, 2021, 15, 20.	2.9	12
15	Whole-Genome Sequencing in Diagnostics of Selected Slovenian Undiagnosed Patients with Rare Disorders. Life, 2021, 11, 205.	2.4	5
16	Discovery of a neuromuscular syndrome caused by biallelic variants in ASCC3. Human Genetics and Genomics Advances, 2021, 2, 100024.	1.7	1
17	Diagnostic yield of exome sequencing in myopathies: Experience of a Slovenian tertiary centre. PLoS ONE, 2021, 16, e0252953.	2.5	7
18	Identification of women at risk for hereditary breast and ovarian cancer in a sample of 1000 Slovenian women: a comparison of guidelines. BMC Cancer, 2021, 21, 665.	2.6	0

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19	A Systematic Review of Parkinson's Disease Pharmacogenomics: Is There Time for Translation into the Clinics?. International Journal of Molecular Sciences, 2021, 22, 7213.	4.1	5
20	MYH7-related disorders in two Bulgarian families: Novel variants in the same region associated with different clinical manifestation and disease penetrance. Neuromuscular Disorders, 2021, 31, 633-641.	0.6	2
21	Disease Progression in CNGA3 and CNGB3 Retinopathy; Characteristics of Slovenian Cohort and Proposed OCT Staging Based on Pooled Data from 126 Patients from 7 Studies. Current Issues in Molecular Biology, 2021, 43, 941-957.	2.4	4
22	Increasing Genomic Literacy Through National Genomic Projects. Frontiers in Genetics, 2021, 12, 693253.	2.3	12
23	Potential protective role of a NOD2 polymorphism in the susceptibility to multiple sclerosis is not associated with interferon therapy. Biomedical Reports, 2021, 15, 100.	2.0	0
24	Editorial: Prognostication of Heart Failure Evolution: From Circulating Biomarkers to Genetic Risk Predictive Score. Frontiers in Cardiovascular Medicine, 2021, 8, 687232.	2.4	1
25	Clinical and Histopathological Features of Gelsolin Amyloidosis Associated with a Novel GSN Variant p.Glu580Lys. International Journal of Molecular Sciences, 2021, 22, 1084.	4.1	8
26	Mucopolysaccharidosis III: Molecular basis and treatment. Pediatric Endocrinology, Diabetes and Metabolism, 2021, 27, 201-208.	0.7	7
27	Children with cavernous malformations of the central nervous system. European Journal of Paediatric Neurology, 2021, 35, 61-66.	1.6	2
28	MicroRNA-Target Interaction Regulatory Network in Alzheimer's Disease. Journal of Personalized Medicine, 2021, 11, 1275.	2.5	11
29	Current State of Compulsory Basic and Clinical Courses in Genetics for Medical Students at Medical Faculties in Balkan Countries With Slavic Languages. Frontiers in Genetics, 2021, 12, 793834.	2.3	0
30	Lack of association between C282Y and H63D polymorphisms in the hemochromatosis gene and risk of multiple sclerosis: A meta‑analysis. Biomedical Reports, 2021, 16, 12.	2.0	2
31	Diagnostic and Clinical Utility of Clinical Exome Sequencing in Children With Moderate and Severe Global Developmental Delay / Intellectual Disability. Journal of Child Neurology, 2020, 35, 116-131.	1.4	22
32	Genetic testing offer for inherited neuromuscular diseases within the EURO-NMD reference network: A European survey study. PLoS ONE, 2020, 15, e0239329.	2.5	6
33	Telegenetics. Current Opinion in Pediatrics, 2020, 32, 739-741.	2.0	5
34	The Role of microRNAs in Heart Failure: A Systematic Review. Frontiers in Cardiovascular Medicine, 2020, 7, 161.	2.4	23
35	Improved Criteria for the Classification of Titin Variants in Inherited Skeletal Myopathies. Journal of Neuromuscular Diseases, 2020, 7, 153-166.	2.6	18
36	Genetics-Related Activities in Everyday Practice of Family Physicians in Slovenia. Public Health Genomics, 2020, 23, 230-236.	1.0	1

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37	Development of an Algorithm for Determining of Genetic Risk at the Primary Healthcare Level – A New Tool for Primary Prevention: A Study Protocol. Zdravstveno Varstvo, 2020, 59, 27-32.	0.9	3
38	DNMT3B rs1569686 and rs2424913 gene polymorphisms are associated with positive family history of preterm birth and smoking status. Croatian Medical Journal, 2020, 61, 8-17.	0.7	8
39	Title is missing!. , 2020, 15, e0239329.		0
40	Title is missing!. , 2020, 15, e0239329.		0
41	Title is missing!. , 2020, 15, e0239329.		0
42	Title is missing!. , 2020, 15, e0239329.		0
43	Actionable Pharmacogenetic Variation in the Slovenian Genomic Database. Frontiers in Pharmacology, 2019, 10, 240.	3.5	10
44	Combination of QFâ€PCR and aCCH is an efficient diagnostic strategy for the detection of chromosome aberrations in recurrent miscarriage. Molecular Genetics & Genomic Medicine, 2019, 7, e980.	1.2	7
45	Vaginal Microbiome Signature Is Associated With Spontaneous Preterm Delivery. Frontiers in Medicine, 2019, 6, 201.	2.6	71
46	Multiple Sclerosis patients carry an increased burden of exceedingly rare genetic variants in the inflammasome regulatory genes. Scientific Reports, 2019, 9, 9171.	3.3	33
47	PEDIA: prioritization of exome data by image analysis. Genetics in Medicine, 2019, 21, 2807-2814.	2.4	58
48	The frequency of CNV s in a cohort population of consecutive fetuses with congenital anomalies after the termination of pregnancy. Molecular Genetics & amp; Genomic Medicine, 2019, 7, e658.	1.2	3
49	Circadian clock genes and myocardial infarction in patients with type 2 diabetes mellitus. Gene, 2019, 701, 98-103.	2.2	19
50	Pharmacogenomics of Multiple Sclerosis: A Systematic Review. Frontiers in Neurology, 2019, 10, 134.	2.4	11
51	Circadian clock genes and circadian phenotypes in patients with myocardial infarction. Advances in Medical Sciences, 2019, 64, 224-229.	2.1	18
52	MiRNA as biomarker for uveitis - A systematic review of the literature. Gene, 2019, 696, 162-175.	2.2	16
53	The role of circadian rhythm in male reproduction. Current Opinion in Endocrinology, Diabetes and Obesity, 2019, 26, 313-316.	2.3	14
54	Rare missense TUBGCP5 gene variant in a patient with primary microcephaly. European Journal of Medical Genetics, 2019, 62, 103598.	1.3	22

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55	MON-220 Androstendione Affects Endometrial Gene Expression Profile During Window of Implantation in Obese Infertile Women with PCOS. Journal of the Endocrine Society, 2019, 3, .	0.2	1
56	Mutations in <i>SCN3A</i> cause early infantile epileptic encephalopathy. Annals of Neurology, 2018, 83, 703-717.	5.3	69
57	Nonsyndromic cleft palate: An association study at GWAS candidate loci in a multiethnic sample. Birth Defects Research, 2018, 110, 871-882.	1.5	11
58	Sarcoidosis Related Novel Candidate Genes Identified by Multi-Omics Integrative Analyses. OMICS A Journal of Integrative Biology, 2018, 22, 322-331.	2.0	10
59	Transcriptomic Biomarkers for Huntington's Disease: Are Gene Expression Signatures in Whole Blood Reliable Biomarkers?. OMICS A Journal of Integrative Biology, 2018, 22, 283-294.	2.0	9
60	Diagnostic exome sequencing of syndromic epilepsy patients in clinical practice. Clinical Genetics, 2018, 93, 1057-1062.	2.0	39
61	Diagnostic efficacy and new variants in isolated and complex autism spectrum disorder using molecular karyotyping. Journal of Applied Genetics, 2018, 59, 179-185.	1.9	12
62	The challenges of the expanded availability of genomic information: an agenda-setting paper. Journal of Community Genetics, 2018, 9, 103-116.	1.2	45
63	Comprehensive use of extended exome analysis improves diagnostic yield in rare disease: a retrospective survey in 1,059 cases. Genetics in Medicine, 2018, 20, 303-312.	2.4	57
64	Association of Circadian Rhythm with Myocardial Infarction. Acta Clinica Croatica, 2018, 57, 480-486.	0.2	4
65	Recessive DNAH9 Loss-of-Function Mutations Cause Laterality Defects and Subtle Respiratory Ciliary-Beating Defects. American Journal of Human Genetics, 2018, 103, 995-1008.	6.2	92
66	Diagnostic Testing in Epilepsy Genetics Clinical Practice. , 2018, , .		0
67	Association between angiotensin-converting enzyme gene insertion/deletion polymorphism and susceptibility to preterm birth: A case-control study and meta-analysis. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2018, 231, 122-128.	1.1	2
68	DNA Methylation Profiles in Whole Blood of Huntington's Disease Patients. Frontiers in Neurology, 2018, 9, 655.	2.4	18
69	Genetic variations in circadian rhythm genes and susceptibility for myocardial infarction. Genetics and Molecular Biology, 2018, 41, 403-409.	1.3	21
70	Microduplication in the 2p16.1p15 chromosomal region linked to developmental delay and intellectual disability. Molecular Cytogenetics, 2018, 11, 39.	0.9	4
71	Genetic variation in the CLOCK gene is associated with idiopathic recurrent spontaneous abortion. PLoS ONE, 2018, 13, e0196345.	2.5	16
72	Diagnostic outcomes of exome sequencing in patients with syndromic or non-syndromic hearing loss. PLoS ONE, 2018, 13, e0188578.	2.5	33

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73	Association of circadian rhythm genes ARNTL/BMAL1 and CLOCK with multiple sclerosis. PLoS ONE, 2018, 13, e0190601.	2.5	34
74	Screening for Rare Genetic Variants Associated with Atherosclerosis: Opportunity for Personalized Medicine. Current Vascular Pharmacology, 2018, 17, 25-28.	1.7	7
75	The lack of association between angiotensinâ€converting enzyme gene insertion/deletion polymorphism and nicotine dependence in multiple sclerosis. Brain and Behavior, 2017, 7, e00600.	2.2	3
76	Angiotensin-converting enzyme insertion/deletion gene polymorphism and interferon-β treatment response in multiple sclerosis patients. Pharmacogenetics and Genomics, 2017, 27, 232-235.	1.5	7
77	Identification of rare genetic variation of NLRP1 gene in familial multiple sclerosis. Scientific Reports, 2017, 7, 3715.	3.3	53
78	Telegenetics: an Update on Availability and Use of Telemedicine in Clinical Genetics Service. Journal of Medical Systems, 2017, 41, 21.	3.6	46
79	Functional single nucleotide polymorphisms of matrix metalloproteinase 7 and 12 genes in idiopathic recurrent spontaneous abortion. Journal of Assisted Reproduction and Genetics, 2017, 34, 365-371.	2.5	2
80	Genetic Interactions in Nonsyndromic Orofacial Clefts in Europe—EUROCRAN Study. Cleft Palate-Craniofacial Journal, 2017, 54, 623-630.	0.9	18
81	De Novo Mutations in SLC25A24 Cause a Disorder Characterized by Early Aging, Bone Dysplasia, Characteristic Face, and Early Demise. American Journal of Human Genetics, 2017, 101, 844-855.	6.2	51
82	Transcriptome Profiling Uncovers Potential Common Mechanisms in Fetal Trisomies 18 and 21. OMICS A Journal of Integrative Biology, 2017, 21, 565-570.	2.0	8
83	A Single Nucleotide Polymorphism of <scp>DNA</scp> methyltransferase 3B gene is a risk factor for recurrent spontaneous abortion. American Journal of Reproductive Immunology, 2017, 78, e12765.	1.2	8
84	Systematic review and meta-analysis of genetic association studies in idiopathic recurrent spontaneous abortion. Fertility and Sterility, 2017, 107, 150-159.e2.	1.0	110
85	Impact of prenatal screening on the prevalence of Down syndrome in Slovenia. PLoS ONE, 2017, 12, e0180348.	2.5	8
86	Genetic variation in leptin and leptin receptor genes is a risk factor for idiopathic recurrent spontaneous abortion. Croatian Medical Journal, 2016, 57, 566-571.	0.7	5
87	Clinical utility of array comparative genomic hybridisation in prenatal setting. BMC Medical Genetics, 2016, 17, 81.	2.1	16
88	Phenotype-driven gene target definition in clinical genome-wide sequencing data interpretation. Genetics in Medicine, 2016, 18, 1102-1110.	2.4	41
89	Angiotensin-converting enzyme insertion/deletion gene polymorphism in multiple sclerosis: a meta-analysis. Neurological Sciences, 2016, 37, 1955-1959.	1.9	4
90	Insertion/deletion polymorphism in intron 16 of ACE gene in idiopathic recurrent spontaneous abortion: case-control study, systematic review and meta-analysis. Reproductive BioMedicine Online, 2016, 32, 237-246.	2.4	14

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91	Reply to Sajantila and Budowle. European Journal of Human Genetics, 2016, 24, 330-330.	2.8	1
92	Responsible implementation of expanded carrier screening. European Journal of Human Genetics, 2016, 24, e1-e12.	2.8	240
93	Epigenetics and Bruxism: Possible Role of Epigenetics in the Etiology of Bruxism. International Journal of Prosthodontics, 2015, 28, 594-599.	1.7	15
94	The influence of cytokine gene polymorphisms on the risk of developing gastric cancer in patients with Helicobacter pylori infection. Radiology and Oncology, 2015, 49, 256-264.	1.7	14
95	Brachytelephalangic chondrodysplasia punctata caused by new small hemizygous deletion in a boy presenting with hearing loss. Molecular Cytogenetics, 2015, 8, 83.	0.9	4
96	Evaluating the SERCA2 and VEGF mRNAs as Potential Molecular Biomarkers of the Onset and Progression in Huntington's Disease. PLoS ONE, 2015, 10, e0125259.	2.5	18
97	Towards a European consensus for reporting incidental findings during clinical NGS testing. European Journal of Human Genetics, 2015, 23, 1601-1606.	2.8	85
98	The â~'2549 insertion/deletion polymorphism in the promoter region of the VEGFA gene in couples with idiopathic recurrent spontaneous abortion. Journal of Assisted Reproduction and Genetics, 2015, 32, 1789-1794.	2.5	9
99	MMP-2 â^ 1575G/A polymorphism modifies the onset of optic neuritis as a first presenting symptom in MS?. Journal of Neuroimmunology, 2015, 286, 13-15.	2.3	8
100	Therapeutic perspectives of epigenetically active nutrients. British Journal of Pharmacology, 2015, 172, 2756-2768.	5.4	99
101	Direct-to-consumer genetic testing in Slovenia: availability, ethical dilemmas and legislation. Biochemia Medica, 2015, 25, 84-89.	2.7	6
102	Cornelia de Lange syndrome caused by heterozygous deletions of chromosome 8q24: Comments on the article by Pereza et al. [2012]. American Journal of Medical Genetics, Part A, 2015, 167, 1426-1427.	1.2	8
103	Non-invasive prenatal testing for aneuploidy and beyond: challenges of responsible innovation in prenatal screening. Summary and recommendations. European Journal of Human Genetics, 2015, , .	2.8	13
104	Non-invasive prenatal testing for aneuploidy and beyond: challenges of responsible innovation in prenatal screening. European Journal of Human Genetics, 2015, 23, 1438-1450.	2.8	260
105	A critical update on endothelial nitric oxide synthase gene variations in women with idiopathic recurrent spontaneous abortion: genetic association study, systematic review and meta-analyses. Molecular Human Reproduction, 2015, 21, 466-478.	2.8	14
106	Family history based approach in risk prediction for Parkinson's disease: Additional contribution of familial associated disorders. Genetika, 2015, 47, 303-310.	0.4	0
107	Omics Technologies and Neovascular Ocular Disorders. BioMed Research International, 2014, 2014, 1-2.	1.9	1
108	Functional Polymorphisms of Matrix Metalloproteinases 1 and 9 Genes in Women with Spontaneous Preterm Birth. Disease Markers, 2014, 2014, 1-7.	1.3	13

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109	The Role of TPA I/D and PAI-1 4G/5G Polymorphisms in Multiple Sclerosis. Disease Markers, 2014, 2014, 1-8.	1.3	10
110	An international effort towards developing standards for best practices in analysis, interpretation and reporting of clinical genome sequencing results in the CLARITY Challenge. Genome Biology, 2014, 15, R53.	9.6	101
111	Recommendations for reporting results of diagnostic genetic testing (biochemical, cytogenetic and) Tj ETQq1 1	0.784314 2.8	4 rgBT /Overloo
112	Strong Association of Variants around <i>FOXE1</i> and Orofacial Clefting. Journal of Dental Research, 2014, 93, 376-381.	5.2	51
113	CSTM1-null and CSTT1-null genotypes are associated with essential arterial hypertension in patients with type 2 diabetes. Clinical Biochemistry, 2014, 47, 574-577.	1.9	17
114	Genetic variation in tissue inhibitors of metalloproteinases as a risk factorÂfor idiopathic recurrent spontaneous abortion. Fertility and Sterility, 2013, 99, 1923-1929.	1.0	7
115	Y chromosome azoospermia factor region microdeletions are not associated with idiopathic recurrent spontaneous abortion in a Slovenian population: association study and literature review. Fertility and Sterility, 2013, 99, 1663-1667.	1.0	9
116	Integration of Data from Omic Studies with the Literature-Based Discovery towards Identification of Novel Treatments for Neovascularization in Diabetic Retinopathy. BioMed Research International, 2013, 2013, 1-7.	1.9	5
117	Analysis of susceptibility loci for nonsyndromic orofacial clefting in a European trio sample. American Journal of Medical Genetics, Part A, 2013, 161, 2545-2549.	1.2	21
118	Correlates of Depression in the Slovenian Working Population. Arhiv Za Higijenu Rada I Toksikologiju, 2013, 64, 489-495.	0.7	7
119	Using Literature-based Discovery to Identify Novel Therapeutic Approaches. Cardiovascular and Hematological Agents in Medicinal Chemistry, 2013, 11, 14-24.	1.0	45
120	Genetic Variation in Circadian Rhythm Genes CLOCK and ARNTL as Risk Factor for Male Infertility. PLoS ONE, 2013, 8, e59220.	2.5	32
121	Family History as a Predictor for Disease Risk in Healthy Individuals: A Cross-Sectional Study in Slovenia. PLoS ONE, 2013, 8, e80333.	2.5	9
122	Expression Signature as a Biomarker for Prenatal Diagnosis of Trisomy 21. PLoS ONE, 2013, 8, e74184.	2.5	27
123	Detection of Thrombophilic Mutations Related to Spontaneous Abortions by a Multiplex SNaPshot Method. Genetic Testing and Molecular Biomarkers, 2012, 16, 259-264.	0.7	12
124	Matrix metalloproteinases 1, 2, 3 and 9 functional single-nucleotide polymorphisms in idiopathic recurrent spontaneous abortion. Reproductive BioMedicine Online, 2012, 24, 567-575.	2.4	33
125	The insulin-like growth factor 2 receptor gene Gly1619Arg polymorphism and idiopathic recurrent spontaneous abortion. Journal of Maternal-Fetal and Neonatal Medicine, 2012, 25, 429-431.	1.5	4
126	Histamine N-methyltransferase Thr105Ile polymorphism is associated with Parkinson's disease. Neurobiology of Aging, 2012, 33, 836.e1-836.e3.	3.1	29

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127	Third case of 8q23.3â€q24.13 deletion in a patient with Langer–Giedion syndrome phenotype without <i>TRPS1</i> gene deletion. American Journal of Medical Genetics, Part A, 2012, 158A, 659-663.	1.2	20
128	Integrative â€~Omic' Approach Towards Understanding the Nature of Human Diseases. Balkan Journal of Medical Genetics, 2012, 15, 45-50.	0.5	5
129	Association between male infertility and genetic variability at the PON1/2 and GSTM1/T1 gene loci. Reproductive BioMedicine Online, 2011, 23, 105-110.	2.4	26
130	A new case of rare proximal 3q13 interstitial deletion. Open Medicine (Poland), 2011, 6, 625-630.	1.3	2
131	Angiotensin-Converting Enzyme Gene Polymorphism in Patients with Multiple Sclerosis from Bosnia and Herzegovina. Genetic Testing and Molecular Biomarkers, 2011, 15, 835-838.	0.7	5
132	Positional integratomic approach in identification of genomic candidate regions for Parkinson's disease. Bioinformatics, 2011, 27, 1971-1978.	4.1	15
133	Low frequency of HFE gene mutations in Croatian patients suspected of having hereditary hemochromatosis. Medical Science Monitor, 2011, 17, CR552-CR556.	1.1	2
134	Autonomic dysfunction in presymptomatic and early symptomatic Huntington's disease. Acta Neurologica Scandinavica, 2010, 121, 392-399.	2.1	49
135	SATB2 haploinsufficiency in patients with cleft palate. Open Medicine (Poland), 2010, 5, 318-321.	1.3	0
136	Rasch-based high-dimensionality data reduction and class prediction with applications to microarray gene expression data. Expert Systems With Applications, 2010, 37, 5178-5185.	7.6	15
137	Chi-square-based Scoring Function for Categorization of MEDLINE Citations. Methods of Information in Medicine, 2010, 49, 371-378.	1.2	13
138	Interleukin 7 receptor alpha polymorphism rs6897932 and susceptibility to multiple sclerosis in the Western Balkans. Multiple Sclerosis Journal, 2010, 16, 533-536.	3.0	11
139	Combining Semantic Relations and DNA Microarray Data for Novel Hypotheses Generation. Lecture Notes in Computer Science, 2010, , 53-61.	1.3	15
140	Lack of association of immune-response-gene polymorphisms with susceptibility to sarcoidosis in Slovenian patients. Genetics and Molecular Research, 2010, 9, 58-68.	0.2	2
141	Interstitial deletion 2p11.2–p12: Further delineation. American Journal of Medical Genetics, Part A, 2009, 149A, 2324-2326.	1.2	7
142	Gene expression changes in blood as a putative biomarker for Huntington's disease. Movement Disorders, 2009, 24, 2277-2281.	3.9	28
143	Epidemiology of Huntington's disease in Slovenia. Acta Neurologica Scandinavica, 2009, 119, 371-375.	2.1	18
144	Polymorphisms in genes involved in folate metabolism as maternal risk factors for Down syndrome — meta-analysis. Open Medicine (Poland), 2009, 4, 395-408.	1.3	6

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145	Association between genetic polymorphisms in cytokine genes and recurrent miscarriage – a meta-analysis. Reproductive BioMedicine Online, 2009, 19, 406-414.	2.4	24
146	Semantic relations for interpreting DNA microarray data. AMIA Annual Symposium proceedings, 2009, 255-9.	0.2	2
147	Search for sarcoidosis candidate genes by integration of data from genomic, transcriptomic and proteomic studies. Medical Science Monitor, 2009, 15, SR22-8.	1.1	15
148	Distribution of HFE gene mutations in Slovenian patients with hereditary hemochromatosis. Annals of Hematology, 2008, 87, 667-669.	1.8	2
149	Genital anomalies in a patient with Treacher Collins syndrome. American Journal of Medical Genetics, Part A, 2008, 146A, 2169-2171.	1.2	3
150	Mutations in C2orf37, Encoding a Nucleolar Protein, Cause Hypogonadism, Alopecia, Diabetes Mellitus, Mental Retardation, and Extrapyramidal Syndrome. American Journal of Human Genetics, 2008, 83, 684-691.	6.2	121
151	ORIGINAL ARTICLE: Genetic Predisposition to Idiopathic Recurrent Spontaneous Abortion: Contribution of Genetic Variations in IGFâ€2 and H19 Imprinted Genes. American Journal of Reproductive Immunology, 2008, 60, 111-117.	1.2	24
152	Meta analysis: Haplotype or single polymorphism analysis. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2008, 136, 260-261.	1.1	0
153	Incidence of the 35delC/CJB2 mutation in low-risk newborns. Journal of Maternal-Fetal and Neonatal Medicine, 2008, 21, 463-468.	1.5	5
154	Heart-hand syndrome of Slovenian type: a new kind of laminopathy. Journal of Medical Genetics, 2008, 45, 666-671.	3.2	47
155	PAI and TPA gene polymorphisms in multiple sclerosis. Multiple Sclerosis Journal, 2008, 14, 243-247.	3.0	13
156	Local and genetic determinants of vascular endothelial growth factor expression in advanced proliferative diabetic retinopathy. Molecular Vision, 2008, 14, 1382-7.	1.1	54
157	Tumor Necrosis Factor-α-308 Gene Polymorphism in Croatian and Slovenian Multiple Sclerosis Patients. European Neurology, 2007, 57, 203-207.	1.4	22
158	Genetic polymorphisms in vasoactive genes and preeclampsia: A meta-analysis. European Journal of Obstetrics, Gynecology and Reproductive Biology, 2007, 131, 115-126.	1.1	69
159	Epidemiology of multiple sclerosis in western Herzegovina. Clinical Neurology and Neurosurgery, 2007, 109, 779-783.	1.4	15
160	Association of Vascular Endothelial Growth Factor Gene Polymorphism with Myocardial Infarction in Patients with Type 2 Diabetes. Cardiology, 2007, 107, 291-295.	1.4	53
161	Polymorphisms in the Interleukin-12/18 Genes and Recurrent Spontaneous Abortion. American Journal of Reproductive Immunology, 2007, 58, 403-408.	1.2	20
162	Analysis of the CAG repeat number in exon 1 of the androgen receptor gene in Slovene men with idiopathic azoospermia and oligoasthenoteratozoospermia. Asian Journal of Andrology, 2007, 9, 280-282.	1.6	7

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163	Patients with primary cataract as a genetic pool of DMPK protomutation. Journal of Human Genetics, 2007, 52, 123-128.	2.3	17
164	Role of genetic polymorphisms in ACE and TNF-α gene in sarcoidosis: a meta-analysis. Journal of Human Genetics, 2007, 52, 836-847.	2.3	44
165	Literature Based Discovery Support System and Its Application to Disease Gene Identification. Lecture Notes in Computer Science, 2007, , 307-326.	1.3	3
166	Region with persistent high frequency of multiple sclerosis in Croatia and Slovenia. Journal of the Neurological Sciences, 2006, 247, 169-172.	0.6	21
167	Analysis of the hemochromatosis mutations C282Y and H63D in infertile men. Fertility and Sterility, 2006, 86, 1796-1798.	1.0	6
168	Angiotensin-converting enzyme I/D gene polymorphism and risk of multiple sclerosis. Acta Neurologica Scandinavica, 2006, 114, 374-377.	2.1	20
169	Preliminary analysis of AZFb region duplication by quantitative real-time PCR. Human Reproduction, 2006, 21, 753-754.	0.9	2
170	No association of CCR5D32 gene mutation with multiple sclerosis in Croatian and Slovenian patients. Multiple Sclerosis Journal, 2006, 12, 360-362.	3.0	15
171	Association between the apolipoprotein B signal peptide gene insertion/deletion polymorphism and male infertility. Molecular Human Reproduction, 2006, 12, 777-779.	2.8	15
172	Exploiting semantic relations for literature-based discovery. AMIA Annual Symposium proceedings, 2006, , 349-53.	0.2	59
173	Oestrogen receptor dinucleotide (TA) polymorphism affects total and LDL cholesterol but does not predict premature myocardial infarction in Caucasian men. Acta Cardiologica, 2005, 60, 307-309.	0.9	1
174	Human Y-specific STR haplotypes in the Western Croatian population sample. Forensic Science International, 2005, 149, 257-261.	2.2	17
175	Familial progressive sinoatrial and atrioventricular conduction disease of adult onset with sudden death, dilated cardiomyopathy, and brachydactyly. A new type of heart-hand syndrome?. Clinical Genetics, 2005, 68, 155-160.	2.0	21
176	Y microdeletions in the Istria county, Croatia. Asian Journal of Andrology, 2005, 7, 213-216.	1.6	15
177	Using literature-based discovery to identify disease candidate genes. International Journal of Medical Informatics, 2005, 74, 289-298.	3.3	219
178	Aldose reductase (AC)n gene polymorphism and susceptibility to diabetic retinopathy in Type 2 diabetes in Caucasians. Journal of Diabetes and Its Complications, 2005, 19, 70-73.	2.3	33
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