Mikhail I Voevoda

List of Publications by Year
in descending order

88	7,176	279487 23	66788 78
papers	citations	h-index	g-index

91	91	91	11987
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Associations of APOE Gene Variants rs429358 and rs7412 with Parameters of the Blood Lipid Profile and the Risk of Myocardial Infarction and Death in a White Population of Western Siberia. Current Issues in Molecular Biology, 2022, 44, 1713-1724.	1.0	4
2	The Mutation Spectrum of Maturity Onset Diabetes of the Young (MODY)-Associated Genes among Western Siberia Patients. Journal of Personalized Medicine, 2021, 11, 57.	1.1	12
3	The Risk of Type 2 Diabetes Mellitus in a Russian Population Cohort According to Data from the HAPIEE Project. Journal of Personalized Medicine, 2021, 11, 119.	1.1	9
4	Basic Research in Endocrinology: A Modern Strategy for the Development and Technologies of Personalized Medicine. Journal of Personalized Medicine, 2021, 11, 895.	1.1	0
5	IgG Study of Blood Sera of Patients with COVID-19. Pathogens, 2021, 10, 1421.	1.2	9
6	Analysis of Rare Variants in Genes Related to Lipid Metabolism in Patients with Familial Hypercholesterolemia in Western Siberia (Russia). Journal of Personalized Medicine, 2021, 11, 1232.	1.1	6
7	Impact of Mitochondrial DNA Mutations on Carotid Intima-Media Thickness in the Novosibirsk Region. Life, 2020, 10, 160.	1.1	4
8	Analysis of APPL1 Gene Polymorphisms in Patients with a Phenotype of Maturity Onset Diabetes of the Young. Journal of Personalized Medicine, 2020, 10, 100.	1.1	13
9	Data on association of mitochondrial heteroplasmy with carotid intima-media thickness in subjects from Russian and Kazakh populations. Data in Brief, 2020, 29, 105136.	0.5	7
10	Potential regulatory SNPs in the ATXN7L3B and KRT15 genes are associated with gender-specific colorectal cancer risk. Personalized Medicine, 2020, 17, 43-54.	0.8	5
11	A rare splice site mutation in the gene encoding glucokinase/hexokinase 4 in a patient with MODY type 2. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 24, 299-305.	0.4	1
12	Ethnicity-specific distribution of TRPM8 gene variants in Eurasian populations: signs of selection. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 24, 292-298.	0.4	1
13	The rs78378222 prevalence and the copy loss of the protective allele A in the tumor tissue of diffuse large B-cell lymphoma. PeerJ, 2020, 8, e10335.	0.9	4
14	Effects of smoking on the level of sp-a and sp-d surfactant proteins in the blood of patients without bronchopulmonary diseases. Bulletin of Siberian Medicine, 2020, 19, 104-111.	0.1	2
15	Association between leukocyte telomere length and specific antibody levels after vaccination against tick-borne encephalitis. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 23, 1026-1031.	0.4	1
16	Changes induced in mouse lipid metabolism by simultaneous impact of antisense oligonucleotide derivatives to <i>apoB</i>, <i>PCSK9</i>, and <i>apoCIII</i> mRNAs. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 23, 1020-1025.	0.4	0
17	Analysis of Polymorphism rs1333049 (Located at 9P21.3) in the White Population of Western Siberia and Associations with Clinical and Biochemical Markers. Biomolecules, 2019, 9, 290.	1.8	6
18	Effects of alirocumab on cardiovascular and metabolic outcomes after acute coronary syndrome in patients with or without diabetes: a prespecified analysis of the ODYSSEY OUTCOMES randomised controlled trial. Lancet Diabetes and Endocrinology,the, 2019, 7, 618-628.	5.5	207

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19	Searching for Signatures of Cold Climate Adaptation in TRPM8 Gene in Populations of East Asian Ancestry. Frontiers in Genetics, 2019, 10, 759.	1.1	9
20	SA76SLEEP DISTURBANCES AND THE VNTR POLYMORPHISM OF CANDIDATE GENES DRD4 AND DAT IN MALE POPULATION 25-64 YEARS IN RUSSIA/SIBERIA: WHO PROGRAM MONICA-PSYCHOSOCIAL. European Neuropsychopharmacology, 2019, 29, S1229.	0.3	0
21	Exome-wide search and functional annotation of genes associated in patients with severe tick-borne encephalitis in a Russian population. BMC Medical Genomics, 2019, 12, 61.	0.7	9
22	The association between the FTO gene variant and alcohol consumption and binge and problem drinking in different gene-environment background: The HAPIEE study. Gene, 2019, 707, 30-35.	1.0	8
23	Exome-wide survey of the Siberian Caucasian population. BMC Medical Genetics, 2019, 20, 51.	2.1	2
24	Clinical aspects of TP53 gene inactivation in diffuse large B-cell lymphoma. BMC Medical Genomics, 2019, 12, 35.	0.7	24
25	Association of RS708272 (CETP Gene Variant) with Lipid Profile Parameters and the Risk of Myocardial Infarction in the White Population of Western Siberia. Biomolecules, 2019, 9, 739.	1.8	8
26	Diabetes mellitus associated with the mutation of the ABCC8 gene (MODY 12): features of clinical course and therapy. Diabetes Mellitus, 2019, 22, 88-94.	0.5	2
27	Gender features of respiratory symptoms in the young population. , 2019, , .		0
28	A Case of Maturity Onset Diabetes of the Young (MODY3) in a Family with a Novel HNF1A Gene Mutation in Five Generations. Diabetes Therapy, 2018, 9, 413-420.	1.2	12
29	A matrix metalloproteinase 9 (MMP9) gene single nucleotide polymorphism is associated with predisposition to tick-borne encephalitis virus-induced severe central nervous system disease. Ticks and Tick-borne Diseases, 2018, 9, 763-767.	1.1	24
30	Leukocyte telomere length and risk of coronary heart disease and stroke mortality: prospective evidence from a Russian cohort. Scientific Reports, 2018, 8, 16627.	1.6	18
31	Genes reveal traces of common recent demographic history for most of the Uralic-speaking populations. Genome Biology, 2018, 19, 139.	3.8	67
32	SIRT1 Allele Frequencies in Depressed Patients of European Descent in Russia. Frontiers in Genetics, 2018, 9, 686.	1.1	6
33	Origin and spread of human mitochondrial DNA haplogroup U7. Scientific Reports, 2017, 7, 46044.	1.6	25
34	New variants in the BRCA1 gene in Buryat Mongol breast cancer patients: Report from two families. Cancer Biomarkers, 2017, 18, 291-296.	0.8	4
35	Prevalence of metabolic syndrome in young adults in Russia. Atherosclerosis, 2017, 263, e172.	0.4	3
36	The polymorphism of cholesterol ester transfer protein gene and lipid profile in men with coronary atherosclerosis. Atherosclerosis, 2017, 263, e186.	0.4	0

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37	Proteomic studies of serum in coronary atherosclerosis. Atherosclerosis, 2017, 263, e195.	0.4	0
38	Analysis of the LDLR gene variability in patients with familial hypercholesterolemia in Russia using targeted high throughput resequencing. Atherosclerosis, 2017, 263, e227.	0.4	2
39	The spectrum of mutations in the CEL gene in early onset diabetes patients. Atherosclerosis, 2017, 263, e259-e260.	0.4	0
40	Clinical case: The development of atherosclerosis in a patient 28 years old with 12 mody diabetes. Atherosclerosis, 2017, 263, e260.	0.4	0
41	The prevalence of dyslipidemia in young patients with diabetes mellitus. Atherosclerosis, 2017, 263, e260-e261.	0.4	0
42	Neuropeptide Receptor Gene (NPSR1) Polymorphism And Sleep Disorders. European Neuropsychopharmacology, 2017, 27, S453.	0.3	1
43	Proprotein convertase subtilisin/kexin type 9 (PCSK9) level in patients with familial hypercholesterolemia in Russia. Atherosclerosis, 2017, 263, e195.	0.4	0
44	The frequency of mutations in the ATP7B gene in Russia. Meta Gene, 2017, 13, 173-176.	0.3	2
45	Association of the genetic markers for myocardial infarction with sudden cardiac death. Indian Heart Journal, 2017, 69, S8-S11.	0.2	13
46	Prevalence of diabetes in the adult population of Novosibirsk. Diabetes Mellitus, 2017, 20, 329-334.	0.5	11
47	Association of IL28B and IL10 gene polymorphism with predisposition to tick-borne encephalitis in a Russian population. Ticks and Tick-borne Diseases, 2016, 7, 808-812.	1.1	34
48	ABCC8-Related Maturity-Onset Diabetes of the Young (MODY12): Clinical Features and Treatment Perspective. Diabetes Therapy, 2016, 7, 591-600.	1.2	41
49	Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242.	13.7	360
50	Validation of the Finnish diabetes risk score (FINDRISC) for the Caucasian population of Siberia. Diabetes Mellitus, 2016, 19, 113-118.	0.5	13
51	Stomach-specific Biomarkers (GastroPanel) Can Predict the Development of Gastric Cancer in a Caucasian Population: A Longitudinal Nested Case-Control Study in Siberia. Anticancer Research, 2016, 36, 247-53.	0.5	15
52	Prognostic impact of the <scp>TP</scp>53 rs1625895 polymorphism in <scp>DLBCL</scp> patients. British Journal of Haematology, 2015, 169, 32-35.	1.2	9
53	The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. PLoS Genetics, 2015, 11, e1005068.	1.5	149
54	Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884.	6.0	449

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55	HMG-coenzyme A reductase inhibition, type 2 diabetes, and bodyweight: evidence from genetic analysis and randomised trials. Lancet, The, 2015, 385, 351-361.	6.3	562
56	Effect ofTRPV4gene polymorphisms on osmotic airway hyperresponsiveness in patients with asthma. , 2015, , .		0
57	Associations of Cold Receptor TRPM8 Gene Single Nucleotide Polymorphism with Blood Lipids and Anthropometric Parameters in Russian Population. Bulletin of Experimental Biology and Medicine, 2014, 157, 757-761.	0.3	13
58	Alcohol consumption and cognitive performance: a <scp>M</scp> endelian randomization study. Addiction, 2014, 109, 1462-1471.	1.7	27
59	Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. Nature, 2014, 505, 87-91.	13.7	821
60	Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413.	13.7	1,179
61	Association between alcohol and cardiovascular disease: Mendelian randomisation analysis based on individual participant data. BMJ, The, 2014, 349, g4164-g4164.	3.0	528
62	CETP Gene Polymorphism in the Caucasian Population of West Siberia and in Groups Contrast by Total Serum Cholesterol Levels. Bulletin of Experimental Biology and Medicine, 2014, 157, 364-367.	0.3	8
63	Prediabetes and Diabetes Prevention Initiatives in Siberia, Russia. , 2014, , 431-447.		1
64	BRCA1 gene mutations frequency estimation by allele-specific real-time PCR of pooled genomic DNA samples. Breast, 2013, 22, 532-536.	0.9	7
65	Autosomal and uniparental portraits of the native populations of Sakha (Yakutia): implications for the peopling of Northeast Eurasia. BMC Evolutionary Biology, 2013, 13, 127.	3.2	106
66	Association of single nucleotide polymorphism rs3775291 in the coding region of the TLR3 gene with predisposition to tick-borne encephalitis in a Russian population. Antiviral Research, 2013, 99, 136-138.	1.9	49
67	Susceptibility to hypoxia and breathing control changes after short-term cold exposures. International Journal of Circumpolar Health, 2013, 72, 21574.	0.5	6
68	The interleukin-6 receptor as a target for prevention of coronary heart disease: a mendelian randomisation analysis. Lancet, The, 2012, 379, 1214-1224.	6.3	886
69	Crosstalk Between theFGFR2andTP53Genes in Breast Cancer: Data from an Association Study and Epistatic Interaction Analysis. DNA and Cell Biology, 2012, 31, 306-316.	0.9	24
70	Single nucleotide polymorphism in the promoter region of the CD209 gene is associated with human predisposition to severe forms of tick-borne encephalitis. Antiviral Research, 2012, 93, 64-68.	1.9	79
71	Apolipoprotein E Gene Polymorphism in Men with Coronary Atherosclerosis in Siberia. Bulletin of Experimental Biology and Medicine, 2011, 150, 355-358.	0.3	7
72	CCL5/RANTES Gene Polymorphisms in Slavonic Patients with Myocardial Infarction. Mediators of Inflammation, 2011, 2011, 1-6.	1.4	17

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73	Blood Levels of Inflammatory and Destructive Biomarkers in Coronary Atherosclerosis of Different Severity. Bulletin of Experimental Biology and Medicine, 2010, 149, 587-590.	0.3	5
74	Variability in the 2â€²â€5â€²â€Oligoadenylate Synthetase Gene Cluster Is Associated with Human Predisposition to Tickâ€Borne Encephalitis Virusâ€“Induced Disease. Journal of Infectious Diseases, 2010, 202, 1813-1818.	1.9	87
75	Association of ADRB1 Gene Polymorphism with Atrial Fibrillation. Genetic Testing and Molecular Biomarkers, 2010, 14, 249-253.	0.3	8
76	TP53 mutations and Arg72Pro polymorphism in breast cancers. Cancer Genetics and Cytogenetics, 2009, 192, 93-95.	1.0	13
77	The macrophage migration inhibitory factor (MIF) gene polymorphism in Czech and Russian patients with myocardial infarction. Clinica Chimica Acta, 2009, 402, 199-202.	0.5	25
78	A counter-clockwise northern route of the Y-chromosome haplogroup N from Southeast Asia towards Europe. European Journal of Human Genetics, 2007, 15, 204-211.	1.4	142
79	Beringian Standstill and Spread of Native American Founders. PLoS ONE, 2007, 2, e829.	1.1	499
80	Risk factors for Alzheimer's disease in Russia: a caseâ€“control study. European Journal of Neurology, 2006, 13, 990-995.	1.7	24
81	Genetic status of p53 in stomach cancer: Somatic mutations and polymorphism of codon 72. Bulletin of Experimental Biology and Medicine, 2006, 141, 243-246.	0.3	22
82	The Western and Eastern Roots of the Saamiâ€”the Story of Genetic â€œOutliersâ€•Told by Mitochondrial DNA and Y Chromosomes. American Journal of Human Genetics, 2004, 74, 661-682.	2.6	202
83	Concerted Changes in the Nucleotide Sequences of the Intragenic Promoter Regions of Eukaryotic Genes for tRNAs of All Specificities. Journal of Molecular Evolution, 2003, 57, 520-532.	0.8	5
84	Origin and Diffusion of mtDNA Haplogroup X. American Journal of Human Genetics, 2003, 73, 1178-1190.	2.6	148
85	Distribution of CCR5-delta 32 gene deletion across the Russian part of Eurasia. Human Genetics, 1998, 102, 695-698.	1.8	25
86	Alcohol consumption and flushing response in natives of Chukotka, Siberia.. Journal of Studies on Alcohol and Drugs, 1995, 56, 194-201.	2.4	2
87	Association of Polymorphism Harbored by Tumor Necrosis Factor Alpha Gene and Sex of Calf with Lactation Performance in Cattle. Asian-Australasian Journal of Animal Sciences, 1970, 26, 1379-1387.	2.4	12
88	Studying accelerated cardiovascular ageing in Russian adults through a novel deep-learning ECG biomarker. Wellcome Open Research, 0, 6, 12.	0.9	8