# Elza Kamilevna Khusnutdinova

List of Publications by Year
in descending order



| 279 | 21,337 | 28274 | 12272 |
|---|---|---|---|
| papers | citations | 55 h-index | 133 g-index |

| 302 | 302 | 302 | 28345 |
|---|---|---|---|
| all docs | docs citations | times ranked | citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | A common founder effect of the splice site variant c.-23â€‰+â€‰1Gâ€‰>â€‰A in GJB2 gene causing autosomal recessive deafness 1A (DFNB1A) in Eurasia. Human Genetics, 2022, 141, 697-707. | 3.8 | 4 |
| 2 | The role of the KIBRA and APOE genes in developing spatial abilities in humans. Vavilovskii Zhurnal Genetiki I Selektsii, 2022, 25, 839-846. | 1.1 | 6 |
| 3 | Rare germline copy number variants (CNVs) and breast cancer risk. Communications Biology, 2022, 5, 65. | 4.4 | 6 |
| 4 | Polygenic risk modeling for prediction of epithelial ovarian cancer risk. European Journal of Human Genetics, 2022, 30, 349-362. | 2.8 | 23 |
| 5 | Common variants in breast cancer risk loci predispose to distinct tumor subtypes. Breast Cancer Research, 2022, 24, 2. | 5.0 | 15 |
| 6 | Pathology of Tumors Associated With Pathogenic Germline Variants in 9 Breast Cancer Susceptibility Genes. JAMA Oncology, 2022, 8, e216744. | 7.1 | 51 |
| 7 | The relationship of lamins with epigenetic factors during aging. Vavilovskii Zhurnal Genetiki I Selektsii, 2022, 26, 40-49. | 1.1 | 0 |
| 8 | Breast cancer risks associated with missense variants in breast cancer susceptibility genes. Genome Medicine, 2022, 14, 51. | 8.2 | 19 |
| 9 | Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 217-228. | 2.5 | 12 |
| 10 | Determination of the phylogenetic origins of the ÃrpÃ¡d Dynasty based on Y chromosome sequencing of BÃ©la the Third. European Journal of Human Genetics, 2021, 29, 164-172. | 2.8 | 18 |
| 11 | CYP3A7*1C allele: linking premenopausal oestrone and progesterone levels with risk of hormone receptor-positive breast cancers. British Journal of Cancer, 2021, 124, 842-854. | 6.4 | 5 |
| 12 | Ethylene-Cytokinin Interaction Determines Early Defense Response of Wheat against Stagonospora nodorum Berk.. Biomolecules, 2021, 11, 174. | 4.0 | 19 |
| 13 | A case-only study to identify genetic modifiers of breast cancer risk for BRCA1/BRCA2 mutation carriers. Nature Communications, 2021, 12, 1078. | 12.8 | 19 |
| 14 | Probable Mechanisms of COVID-19 Pathogenesis. KreativnaÃ¢ HirurgiÃ¢ I OnkologiÃ¢, 2021, 10, 302-310. | 0.3 | 0 |
| 15 | Breast Cancer Risk Genes â€" Association Analysis in More than 113,000 Women. New England Journal of Medicine, 2021, 384, 428-439. | 27.0 | 532 |
| 16 | Specific Features of Ovarian Cancer Metastasis. KreativnaÃ¢ HirurgiÃ¢ I OnkologiÃ¢, 2021, 10, 319-329. | 0.3 | 0 |
| 17 | Autosomal recessive cataract (CTRCT18) in the Yakut population isolate of Eastern Siberia: a novel founder variant in the FYCO1 gene. European Journal of Human Genetics, 2021, 29, 965-976. | 2.8 | 9 |
| 18 | Origin and diffusion of human Y chromosome haplogroup J1-M267. Scientific Reports, 2021, 11, 6659. | 3.3 | 26 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | Investigating the role of osteoprotegerin gene polymorphic variants in osteoporosis. Russian Open Medical Journal, 2021, 10, . | 0.3 | 4 |
| 20 | Novel MicroRNA Binding Site SNPs and the Risk of Clear Cell Renal Cell Carcinoma (ccRCC): A Case-Control Study. Current Cancer Drug Targets, 2021, 21, 203-212. | 1.6 | 6 |
| 21 | Association of Gasdermin B Gene GSDMB Polymorphisms with Risk of Allergic Diseases. Biochemical Genetics, 2021, 59, 1527-1543. | 1.7 | 4 |
| 22 | Neurofibromatosis type 1: results of our own study (Republic of Bashkortostan). Uspehi Molekularnoj Onkologii, 2021, 8, 17-25. | 0.3 | 2 |
| 23 | Pleiotropy-guided transcriptome imputation from normal and tumor tissues identifies candidate susceptibility genes for breast and ovarian cancer. Human Genetics and Genomics Advances, 2021, 2, 100042. | 1.7 | 6 |
| 24 | Functional annotation of the 2q35 breast cancer risk locus implicates a structural variant in influencing activity of a long-range enhancer element. American Journal of Human Genetics, 2021, 108, 1190-1203. | 6.2 | 6 |
| 25 | Mendelian randomisation study of smoking exposure in relation to breast cancer risk. British Journal of Cancer, 2021, 125, 1135-1145. | 6.4 | 9 |
| 26 | The association study of polymorphic variants of hypothalamic-pituitary-adrenal system genes (AVPR1B, OXTR) and aggressive behavior manifestation: a focus on social environment. Research Results in Biomedicine, 2021, 7, 232-244. | 0.5 | 2 |
| 27 | Structural Variability, Expression Profile, and Pharmacogenetic Properties of TMPRSS2 Gene as a Potential Target for COVID-19 Therapy. Genes, 2021, 12, 19. | 2.4 | 22 |
| 28 | Germline variants and breast cancer survival in patients with distant metastases at primary breast cancer diagnosis. Scientific Reports, 2021, 11, 19787. | 3.3 | 2 |
| 29 | Association analysis of serotonin system gene polymorphisms (5-HTT, HTR1B, HTR2A, HTR2C, and TPH1) with the risk of Parkinson's disease in Tatars. Ã‚kutskij Medicinskij Å¾urnal, 2021, , 14-19. | 0.1 | 0 |
| 30 | Analysis of association between histamine receptor gene <i>HRH1</i>, <i>HRH2</i>, <i>HRH3</i>, <i>HRH4</i> polymorphisms and asthma in children. Pulmonologiya, 2021, 31, 729-738. | 0.8 | 0 |
| 31 | Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73. | 21.4 | 120 |
| 32 | Exome sequencing study of Russian breast cancer patients suggests a predisposing role for USP39. Breast Cancer Research and Treatment, 2020, 179, 731-742. | 2.5 | 9 |
| 33 | Breast Cancer Polygenic Risk Score and Contralateral Breast Cancer Risk. American Journal of Human Genetics, 2020, 107, 837-848. | 6.2 | 39 |
| 34 | Genome-wide association study identifies 32 novel breast cancer susceptibility loci from overall and subtype-specific analyses. Nature Genetics, 2020, 52, 572-581. | 21.4 | 265 |
| 35 | Germline HOXB13 mutations p.G84E and p.R217C do not confer an increased breast cancer risk. Scientific Reports, 2020, 10, 9688. | 3.3 | 2 |
| 36 | Transcriptomeâ€wide association study of breast cancer risk by estrogenâ€receptor status. Genetic Epidemiology, 2020, 44, 442-468. | 1.3 | 32 |

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|---|---|---|---|
| 37 | Longitudinal genetic studies of cognitive characteristics. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 24, 87-95. | 1.1 | 10 |
| 38 | Association analysis of amine oxidase 1 AOC1 and histamine-N-methyl-transferase HNMT genes polymorphism with the development of asthma in children. Ã‚kutskij Medicinskij Å¾urnal, 2020, , 20-22. | 0.1 | 0 |
| 39 | Outlook for Neurofi bromatosis Type I Research in the Republic of Bashkortostan. KreativnaÃ¢ HirurgiÃ¢ I OnkologiÃ¢, 2020, 10, 115-121. | 0.3 | 0 |
| 40 | Molecular-genetic bases of development of acute pancreatitis. I P Pavlov Russian Medical Biological Herald, 2020, 28, 536-547. | 0.5 | 2 |
| 41 | Involvement of transposable elements in neurogenesis. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 24, 209-218. | 1.1 | 9 |
| 42 | The role of polymorphic variants of arginase genes (<i>ARG1, ARG2</i>) involved in beta-2-agonist metabolism in the development and course of asthma. Vavilovskii Zhurnal Genetiki I Selektsii, 2020, 24, 391-398. | 1.1 | 3 |
| 43 | A new approach to estimating the prevalence of hereditary hearing loss: An analysis of the distribution of sign language users based on census data in Russia. PLoS ONE, 2020, 15, e0242219. | 2.5 | 1 |
| 44 | Shifts in the Genetic Landscape of the Western Eurasian Steppe Associated with the Beginning and End of the Scythian Dominance. Current Biology, 2019, 29, 2430-2441.e10. | 3.9 | 44 |
| 45 | A Splice Site Variant of CDK12 and Breast Cancer in Three Eurasian Populations. Frontiers in Oncology, 2019, 9, 493. | 2.8 | 4 |
| 46 | The FANCM:p.Arg658* truncating variant is associated with risk of triple-negative breast cancer. Npj Breast Cancer, 2019, 5, 38. | 5.2 | 28 |
| 47 | Two truncating variants in FANCC and breast cancer risk. Scientific Reports, 2019, 9, 12524. | 3.3 | 5 |
| 48 | Opening up new horizons for psychiatric genetics in the Russian Federation: moving toward a national consortium. Molecular Psychiatry, 2019, 24, 1099-1111. | 7.9 | 11 |
| 49 | Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431. | 12.8 | 88 |
| 50 | Y-chromosomal connection between Hungarians and geographically distant populations of the Ural Mountain region and West Siberia. Scientific Reports, 2019, 9, 7786. | 3.3 | 12 |
| 51 | A rare case of Waardenburg syndrome with unilateral hearing loss caused by nonsense variant c.772C>T (p.Arg259*) in the MITF gene in Yakut patient from the Eastern Siberia (Sakha) Tj ETQq1 1 0.784314 rgBT /Overlock | 1.1 | |
| 52 | Comparison of PredictiveIn SilicoTools on Missense Variants inGJB2,GJB6, andGJB3Genes Associated with Autosomal Recessive Deafness 1A (DFNB1A). Scientific World Journal, The, 2019, 2019, 1-9. | 2.1 | 26 |
| 53 | Genome-wide association and transcriptome studies identify target genes and risk loci for breast cancer. Nature Communications, 2019, 10, 1741. | 12.8 | 90 |
| 54 | Fine-scale haplotype mapping of MUT, AACS, SLC6A15 and PRKCA genes indicates association with insulin resistance of metabolic syndrome and relationship with branched chain amino acid metabolism or regulation. PLoS ONE, 2019, 14, e0214122. | 2.5 | 12 |

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|---|---|---|---|
| 55 | Y-chromosomal analysis of clan structure of Kalmyks, the only European Mongol people, and their relationship to Oirat-Mongols of Inner Asia. European Journal of Human Genetics, 2019, 27, 1466-1474. | 2.8 | 8 |
| 56 | Genome-wide association study of germline variants and breast cancer-specific mortality. British Journal of Cancer, 2019, 120, 647-657. | 6.4 | 52 |
| 57 | Polygenic Risk Scores for Prediction of Breast Cancer and Breast Cancer Subtypes. American Journal of Human Genetics, 2019, 104, 21-34. | 6.2 | 711 |
| 58 | Genetic basis of depressive disorders. Vavilovskii Zhurnal Genetiki I Selektsii, 2019, 23, 465-472. | 1.1 | 7 |
| 59 | Association of vascular endothelial growth factor B (VEGFÐ') gene polymorphisms with intracranial aneurysms. Vavilovskii Zhurnal Genetiki I Selektsii, 2019, 22, 992-999. | 1.1 | 0 |
| 60 | The role of miRNA genes participating in VHL-HIF1Î± in clear cell renal cell carcinoma. Urology Herald, 2019, 6, 36-41. | 0.4 | 1 |
| 61 | The role of transposable elements in the ecological morphogenesis under the influence of stress. Vavilovskii Zhurnal Genetiki I Selektsii, 2019, 23, 380-389. | 1.1 | 10 |
| 62 | Epigenetics of suicidal behavior. Vavilovskii Zhurnal Genetiki I Selektsii, 2019, 23, 600-607. | 1.1 | 0 |
| 63 | PROGNOSTIC VALUE OF CYP3A5 AND hOCT1 POLYMORPHIC GENE VARIANTS IN PATIENTS WITH CHRONIC MYELOID LEUKEMIA IN THE REPUBLIC OF BASHKORTOSTAN. Gematologiya I Transfuziologiya, 2019, 64, 165-174. | 0.6 | 0 |
| 64 | Multiancestry association study identifies new asthma risk loci that colocalize with immune-cell enhancer marks. Nature Genetics, 2018, 50, 42-53. | 21.4 | 426 |
| 65 | A novel pathogenic variant c.975G>A (p.Trp325*) in the POU3F4 gene in Yakut family (Eastern Siberia,) Tj ETQq1 1 0.784314 rgBT /O<br>2018, 104, 94-97. | 1.0 | 8 |
| 66 | Gene set enrichment analysis and expression pattern exploration implicate an involvement of neurodevelopmental processes in bipolar disorder. Journal of Affective Disorders, 2018, 228, 20-25. | 4.1 | 14 |
| 67 | Genes reveal traces of common recent demographic history for most of the Uralic-speaking populations. Genome Biology, 2018, 19, 139. | 8.8 | 67 |
| 68 | Branched-Chain Amino Acid Database Integrated in MEDIPAD Software as a Tool for Nutritional Investigation of Mediterranean Populations. Nutrients, 2018, 10, 1392. | 4.1 | 5 |
| 69 | A transcriptome-wide association study of 229,000 women identifies new candidate susceptibility genes for breast cancer. Nature Genetics, 2018, 50, 968-978. | 21.4 | 184 |
| 70 | Recent advances in genetics of aggressive behavior. Vavilovskii Zhurnal Genetiki I Selektsii, 2018, 22, 716-725. | 1.1 | 3 |
| 71 | The role of VNTR aggrecan gene polymorphism in the development of osteoarthritis in women. Vavilovskii Zhurnal Genetiki I Selektsii, 2018, 22, 865-872. | 1.1 | 1 |
| 72 | Identification of alterations in the nucleotide sequence of the chromatin remodeling gene PBRM1 in clear cell renal cell carcinoma patients. Vavilovskii Zhurnal Genetiki I Selektsii, 2018, 22, 873-877. | 1.1 | 0 |

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|---|---|---|---|
| 73 | Assessment of an APOBEC3B truncating mutation, c.783delG, in patients with breast cancer. Breast Cancer Research and Treatment, 2017, 162, 31-37. | 2.5 | 5 |
| 74 | Origin and spread of human mitochondrial DNA haplogroup U7. Scientific Reports, 2017, 7, 46044. | 3.3 | 25 |
| 75 | Identification of 12 new susceptibility loci for different histotypes of epithelial ovarian cancer. Nature Genetics, 2017, 49, 680-691. | 21.4 | 356 |
| 76 | Opinions of hearing parents about the causes of hearing impairment of their children with biallelic GJB2 mutations. Journal of Community Genetics, 2017, 8, 167-171. | 1.2 | 2 |
| 77 | Association analysis identifies 65 new breast cancer risk loci. Nature, 2017, 551, 92-94. | 27.8 | 1,099 |
| 78 | Identification of ten variants associated with risk of estrogen-receptor-negative breast cancer. Nature Genetics, 2017, 49, 1767-1778. | 21.4 | 289 |
| 79 | Genetic aspects of keratoconus development. Russian Journal of Genetics, 2017, 53, 519-527. | 0.6 | 0 |
| 80 | Genetic modifiers of CHEK2*1100delC-associated breast cancer risk. Genetics in Medicine, 2017, 19, 599-603. | 2.4 | 67 |
| 81 | Identification of shared risk loci and pathways for bipolar disorder and schizophrenia. PLoS ONE, 2017, 12, e0171595. | 2.5 | 77 |
| 82 | Between Lake Baikal and the Baltic Sea: genomic history of the gateway to Europe. BMC Genetics, 2017, 18, 110. | 2.7 | 34 |
| 83 | A European Spectrum of Pharmacogenomic Biomarkers: Implications for Clinical Pharmacogenomics. PLoS ONE, 2016, 11, e0162866. | 2.5 | 96 |
| 84 | The study of filaggrin gene mutations and copy number variation in atopic dermatitis patients from Volga-Ural region of Russia. Gene, 2016, 591, 85-89. | 2.2 | 8 |
| 85 | Human Y Chromosome Haplogroup N: A Non-trivial Time-Resolved Phylogeography that Cuts across Language Families. American Journal of Human Genetics, 2016, 99, 163-173. | 6.2 | 98 |
| 86 | The search for new candidate genes involved in ovarian cancer pathogenesis by exome sequencing. Russian Journal of Genetics, 2016, 52, 1105-1109. | 0.6 | 2 |
| 87 | Identification of independent association signals and putative functional variants for breast cancer risk through fine-scale mapping of the 12p11 locus. Breast Cancer Research, 2016, 18, 64. | 5.0 | 31 |
| 88 | Molecular Genetic Investigations of Personality: From Candidate Genes to Genome-wide Associations. , 2016, , 130-154. | | 1 |
| 89 | Mutational landscape of prostate tumors revealed by whole-exome sequencing. Russian Journal of Genetics, 2016, 52, 999-1003. | 0.6 | 1 |
| 90 | The Role of Genetic and Environmental Risk Factors in Aetiology of Suicidal Behaviour. , 2016, , 205-230. | | 0 |

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| 91 | The Simons Genome Diversity Project: 300 genomes from 142 diverse populations. Nature, 2016, 538, 201-206. | 27.8 | 1,216 |
| 92 | Genomic analyses inform on migration events during the peopling of Eurasia. Nature, 2016, 538, 238-242. | 27.8 | 360 |
| 93 | Two novel mutations in gene SPG4 in patients with autosomal dominant spastic paraplegia. Russian Journal of Genetics, 2016, 52, 603-607. | 0.6 | 3 |
| 94 | Genome-Wide Meta-Analyses of Breast, Ovarian, and Prostate Cancer Association Studies Identify Multiple New Susceptibility Loci Shared by at Least Two Cancer Types. Cancer Discovery, 2016, 6, 1052-1067. | 9.4 | 157 |
| 95 | East Eurasian ancestry in the middle of Europe: genetic footprints of Steppe nomads in the genomes of Belarusian Lipka Tatars. Scientific Reports, 2016, 6, 30197. | 3.3 | 14 |
| 96 | Age- and Tumor Subtypeâ€“Specific Breast Cancer Risk Estimates for CHEK2*1100delC Carriers. Journal of Clinical Oncology, 2016, 34, 2750-2760. | 1.6 | 152 |
| 97 | Role of allelic genes of matrix metalloproteinases and their tissue inhibitors in the risk of peptic ulcer disease development. Russian Journal of Genetics, 2016, 52, 320-330. | 0.6 | 5 |
| 98 | Identification of a new locus at 16q12 associated with time to asthma onset. Journal of Allergy and Clinical Immunology, 2016, 138, 1071-1080. | 2.9 | 25 |
| 99 | nZ,(nÂ+Â4)Z-Dienoic fatty acids: a new method for the synthesis and inhibitory action on topoisomerase I and IIÎ±. Medicinal Chemistry Research, 2016, 25, 30-39. | 2.4 | 33 |
| 100 | Spectrum and Frequency of the GJB2 Gene Pathogenic Variants in a Large Cohort of Patients with Hearing Impairment Living in a Subarctic Region of Russia (the Sakha Republic). PLoS ONE, 2016, 11, e0156300. | 2.5 | 21 |
| 101 | Diversity of Y-chromosomal and mtDNA Markers Included in Mediscope Chip within Two Albanian Subpopulations from Croatia and Kosovo: Preliminary Data. Collegium Antropologicum, 2016, 40, 195-8. | 0.2 | 4 |
| 102 | The role of neurotrophins and neurexins genes in the risk of paranoid schizophrenia in Russians and Tatars. Russian Journal of Genetics, 2015, 51, 683-694. | 0.6 | 5 |
| 103 | Association of MUC19 gene polymorphic variants with asthma in Russians based on genome-wide study results. Russian Journal of Genetics, 2015, 51, 1135-1143. | 0.6 | 4 |
| 104 | Genetic Heritage of the Balto-Slavic Speaking Populations: A Synthesis of Autosomal, Mitochondrial and Y-Chromosomal Data. PLoS ONE, 2015, 10, e0135820. | 2.5 | 91 |
| 105 | Prediction of Breast Cancer Risk Based on Profiling With Common Genetic Variants. Journal of the National Cancer Institute, 2015, 107, . | 6.3 | 428 |
| 106 | Polymorphism of brain neurotransmitter system genes: Search for pharmacogenetic markers of haloperidol efficiency in Russians and Tatars. Molecular Biology, 2015, 49, 858-866. | 1.3 | 7 |
| 107 | New genetic loci link adipose and insulin biology to body fat distribution. Nature, 2015, 518, 187-196. | 27.8 | 1,328 |
| 108 | Genetic characterization of populations of the Volga-Ural region according to the variability of the Y-chromosome. Russian Journal of Genetics, 2015, 51, 108-115. | 0.6 | 5 |

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| 109 | The Genetic Legacy of the Expansion of Turkic-Speaking Nomads across Eurasia. PLoS Genetics, 2015, 11, e1005068. | 3.5 | 149 |
| 110 | Global diversity, population stratification, and selection of human copy-number variation. Science, 2015, 349, aab3761. | 12.6 | 293 |
| 111 | Genomic evidence for the Pleistocene and recent population history of Native Americans. Science, 2015, 349, aab3884. | 12.6 | 449 |
| 112 | Current state of research in ethnogenomics: Genome-wide analysis and uniparental markers. Russian Journal of Genetics, 2015, 51, 418-429. | 0.6 | 0 |
| 113 | Genetic and environmental aspects of mathematical disabilities. Russian Journal of Genetics, 2015, 51, 223-230. | 0.6 | 5 |
| 114 | A recent bottleneck of Y chromosome diversity coincides with a global change in culture. Genome Research, 2015, 25, 459-466. | 5.5 | 348 |
| 115 | Stereoselective synthesis of 11-phenylundeca-5Z,9Z-dienoic acid and investigation of its human topoisomerase I and IIÎ± inhibitory activity. Bioorganic and Medicinal Chemistry Letters, 2015, 25, 2405-2408. | 2.2 | 35 |
| 116 | The role of miRNAs in the development of prostate cancer. Russian Journal of Genetics, 2015, 51, 627-641. | 0.6 | 2 |
| 117 | Genetic association of ADRA2A and ADRB3 genes with metabolic syndrome among the Tatars. Russian Journal of Genetics, 2015, 51, 711-714. | 0.6 | 2 |
| 118 | Genome-wide analysis implicates microRNAs and their target genes in the development of bipolar disorder. Translational Psychiatry, 2015, 5, e678-e678. | 4.8 | 67 |
| 119 | Prevalence of the BLM nonsense mutation, p.Q548X, in ovarian cancer patients from Central and Eastern Europe. Familial Cancer, 2015, 14, 145-149. | 1.9 | 12 |
| 120 | The phylogenetic and geographic structure of Y-chromosome haplogroup R1a. European Journal of Human Genetics, 2015, 23, 124-131. | 2.8 | 122 |
| 121 | Brain derived neurotrophic factor gene (BDNF) and personality traits: The modifying effect of season of birth and sex. Progress in Neuro-Psychopharmacology and Biological Psychiatry, 2015, 56, 58-65. | 4.8 | 11 |
| 122 | 11-Phenylundeca-5Z,9Z-dienoic Acid: Stereoselective Synthesis and Dual Topoisomerase I/IIÎ± Inhibition. Current Cancer Drug Targets, 2015, 15, 504-510. | 1.6 | 14 |
| 123 | Mutation Analysis of the ERCC4/FANCQ Gene in Hereditary Breast Cancer. PLoS ONE, 2014, 9, e85334. | 2.5 | 16 |
| 124 | Bioethical issues of preventing hereditary diseases with late onset in the Sakha Republic (Yakutia). International Journal of Circumpolar Health, 2014, 73, 25062. | 1.2 | 8 |
| 125 | Length polymorphism of the B2-VNTR minisatellite repeat of the bradykinin B2 receptor gene in healthy Russians and patients with coronary heart disease. Molecular Biology, 2014, 48, 655-663. | 1.3 | 1 |
| 126 | Association of cytokine gene polymorphisms in peptic ulcer development in the Bashkortostan Republic. Russian Journal of Genetics, 2014, 50, 1316-1325. | 0.6 | 2 |

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|---|---|---|---|
| 127 | Synthesis and Cytotoxicity of Triterpene A-seco-Acid Propargylamides. Chemistry of Natural Compounds, 2014, 50, 853-856. | 0.8 | 7 |
| 128 | Polymorphism of the glutamate receptor genes and risk of paranoid schizophrenia in Russians and Tatars from the Republic of Bashkortostan. Molecular Biology, 2014, 48, 671-680. | 1.3 | 6 |
| 129 | Oxidation of Methyl 2-Cyano-3,4-seco-4(23)-Ene-Ursolate by Ozone. Chemistry of Natural Compounds, 2014, 50, 1037-1041. | 0.8 | 2 |
| 130 | Frequency of CHEK2 gene mutations in breast cancer patients from Republic of Bashkortostan. Molecular Biology, 2014, 48, 46-51. | 1.3 | 7 |
| 131 | Genome-wide association study reveals two new risk loci for bipolar disorder. Nature Communications, 2014, 5, 3339. | 12.8 | 294 |
| 132 | Association of polymorphisms in the toll-like receptor genes with atopic dermatitis in the Republic of Bashkortostan. Molecular Biology, 2014, 48, 227-237. | 1.3 | 2 |
| 133 | Prevalence of PALB2 mutation c.509_510delGA in unselected breast cancer patients from Central and Eastern Europe. Familial Cancer, 2014, 13, 137-142. | 1.9 | 20 |
| 134 | Upper Palaeolithic Siberian genome reveals dual ancestry of Native Americans. Nature, 2014, 505, 87-91. | 27.8 | 821 |
| 135 | Associations between Vitamin D-Binding Protein (DBP) Gene Polymorphism (TAAA)n and Development of Osteoporosis in the Volga-Ural Region of Russia. Bulletin of Experimental Biology and Medicine, 2014, 157, 253-257. | 0.8 | 3 |
| 136 | Association of SLC18A1, TPH1, and RELN gene polymorphisms with risk of paranoid schizophrenia. Molecular Biology, 2014, 48, 546-555. | 1.3 | 16 |
| 137 | Ancient human genomes suggest three ancestral populations for present-day Europeans. Nature, 2014, 513, 409-413. | 27.8 | 1,179 |
| 138 | Arginine-vasopressin receptor gene (AVPR1A, AVPR1B) polymorphisms and their relation to personality traits. Russian Journal of Genetics, 2014, 50, 298-307. | 0.6 | 5 |
| 139 | DNA repair XPCC1 and XPD genes polymorphism as associated with the development of bladder cancer and renal cell carcinoma. Russian Journal of Genetics, 2014, 50, 421-429. | 0.6 | 7 |
| 140 | Examination of structural changes in the transforming growth factor Î² receptor 1 (TGFÎ²R1) gene in patients with chronic heart failure. Russian Journal of Genetics, 2014, 50, 536-542. | 0.6 | 0 |
| 141 | The genetic prehistory of the New World Arctic. Science, 2014, 345, 1255832. | 12.6 | 264 |
| 142 | Age-Related Hearing Impairment (ARHI) Associated with GJB2 Single Mutation IVS1+1G>A in the Yakut Population Isolate in Eastern Siberia. PLoS ONE, 2014, 9, e100848. | 2.5 | 4 |
| 143 | The facile synthesis of the 5Z,9Z-dienoic acids and their topoisomerase I inhibitory activity. Chemical Communications, 2013, 49, 8401. | 4.1 | 53 |
| 144 | MFN2 gene analysis in patients with hereditary motor and sensory neuropathy from Bashkortostan Republic. Russian Journal of Genetics, 2013, 49, 771-777. | 0.6 | 0 |

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|---|---|---|---|
| 145 | Association of YWHAE gene polymorphism with suicidal behavior. Russian Journal of Genetics, 2013, 49, 667-672. | 0.6 | 2 |
| 146 | Analysis of H63D mutation in hemochromatosis (HFE) gene in populations of Central Eurasia. Russian Journal of Genetics, 2013, 49, 237-245. | 0.6 | 1 |
| 147 | Association polymorphic variants of GRIN2B gene with paranoid schizophrenia and response to typical neuroleptics in Russians and Tatars from Bashkortostan Republic. Russian Journal of Genetics, 2013, 49, 962-968. | 0.6 | 11 |
| 148 | Polymorphism of RGS2 gene as genetic marker of schizophrenia risk and pharmacogenetic markers of the efficiency of typical neuroleptics. Molecular Biology, 2013, 47, 814-820. | 1.3 | 13 |
| 149 | Nonsense mutation p.Q548X in BLM, the gene mutated in Bloomâ€™s syndrome, is associated with breast cancer in Slavic populations. Breast Cancer Research and Treatment, 2013, 137, 533-539. | 2.5 | 56 |
| 150 | The role of inflammatory chemokines in lymphoid neoorganogenesis in breast cancer. Biomedicine and Pharmacotherapy, 2013, 67, 363-366. | 5.6 | 28 |
| 151 | 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 |
| 152 | No Evidence from Genome-Wide Data of a Khazar Origin for the Ashkenazi Jews. Human Biology, 2013, 85, 859-900. | 0.2 | 68 |
| 153 | No Evidence from Genome-wide Data of a Khazar Origin fo the Ashkenazi Jews. Human Biology, 2013, 85, 859. | 0.2 | 30 |
| 154 | M06â€…Epidemiology of Huntington's disease in the Republic of Bashkortostan. Journal of Neurology, Neurosurgery and Psychiatry, 2012, 83, A48.2-A48. | 1.9 | 1 |
| 155 | Genome-wide association analysis identifies three new breast cancer susceptibility loci. Nature Genetics, 2012, 44, 312-318. | 21.4 | 256 |
| 156 | Genome-wide meta-analysis identifies 56 bone mineral density loci and reveals 14 loci associated with risk of fracture. Nature Genetics, 2012, 44, 491-501. | 21.4 | 1,100 |
| 157 | The Caucasus as an Asymmetric Semipermeable Barrier to Ancient Human Migrations. Molecular Biology and Evolution, 2012, 29, 1891-1891. | 8.9 | 2 |
| 158 | 9q31.2-rs865686 as a Susceptibility Locus for Estrogen Receptor-Positive Breast Cancer: Evidence from the Breast Cancer Association Consortium. Cancer Epidemiology Biomarkers and Prevention, 2012, 21, 1783-1791. | 2.5 | 17 |
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