Sonja Pavlovic

List of Publications by Year
in descending order

125	1,994	331670	345221
papers	citations	21 h-index	36 g-index

126	126	126	3622
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Systematic documentation and analysis of human genetic variation in hemoglobinopathies using the microattribution approach. Nature Genetics, 2011, 43, 295-301.	21.4	142
2	Frequent deletions of JARID2 in leukemic transformation of chronic myeloid malignancies. American Journal of Hematology, 2012, 87, 245-250.	4.1	107
3	A European Spectrum of Pharmacogenomic Biomarkers: Implications for Clinical Pharmacogenomics. PLoS ONE, 2016, 11, e0162866.	2.5	96
4	Intraâ€articular injection of autologous adiposeâ€derived mesenchymal stem cells in the treatment of knee osteoarthritis. Journal of Gene Medicine, 2018, 20, e3002.	2.8	74
5	Maritime route of colonization of Europe. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 9211-9216.	7.1	71
6	Clinical significance of genetic aberrations in secondary acute myeloid leukemia. American Journal of Hematology, 2012, 87, 1010-1016.	4.1	67
7	Identification and validation of differentially expressed transcripts by RNA-sequencing of formalin-fixed, paraffin-embedded (FFPE) lung tissue from patients with Idiopathic Pulmonary Fibrosis. BMC Pulmonary Medicine, 2017, 17, 15.	2.0	63
8	Association of Bax Expression and Bcl2/Bax Ratio with Clinical and Molecular Prognostic Markers in Chronic Lymphocytic Leukemia. Journal of Medical Biochemistry, 2016, 35, 150-157.	1.7	46
9	Functional analysis of the role of the TPMT gene promoter VNTR polymorphism in TPMT gene transcription. Pharmacogenomics, 2010, 11, 547-557.	1.3	40
10	Epidermal Growth Factor Receptor Gene in Non-Small-Cell Lung Cancer: The Importance of Promoter Polymorphism Investigation. Analytical Cellular Pathology, 2018, 2018, 1-9.	1.4	35
11	Analysis of Thiopurine S-methyltransferase Polymorphism in the Population of Serbia and Montenegro and Mercaptopurine Therapy Tolerance in Childhood Acute Lymphoblastic Leukemia. Therapeutic Drug Monitoring, 2006, 28, 800-806.	2.0	33
12	A gene expression profile associated with relapse of cytogenetically normal acute myeloid leukemia is enriched for leukemia stem cell genes. Leukemia and Lymphoma, 2015, 56, 1126-1128.	1.3	33
13	Importance of early detection and follow-up of FLT3 mutations in patients with acute myeloid leukemia. Annals of Hematology, 2007, 86, 741-747.	1.8	31
14	-174G/C interleukin-6 gene promoter polymorphism predicts therapeutic response to etanercept in rheumatoid arthritis. Rheumatology International, 2013, 33, 1481-1486.	3.0	30
15	Gene Mutation Profiles in Primary Diffuse Large B Cell Lymphoma of Central Nervous System: Next Generation Sequencing Analyses. International Journal of Molecular Sciences, 2016, 17, 683.	4.1	29
16	Pharmacogenomic and Pharmacotranscriptomic Profiling of Childhood Acute Lymphoblastic Leukemia: Paving the Way to Personalized Treatment. Genes, 2019, 10, 191.	2.4	29
17	Use of Wilms Tumor 1 Gene Expression as a Reliable Marker for Prognosis and Minimal Residual Disease Monitoring in Acute Myeloid Leukemia With Normal Karyotype Patients. Clinical Lymphoma, Myeloma and Leukemia, 2017, 17, 312-319.	0.4	26
18	Genomic variation in the MAP3K5 gene is associated with Î²-thalassemia disease severity and hydroxyurea treatment efficacy. Pharmacogenomics, 2013, 14, 469-483.	1.3	25

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19	Genomic profiling supports the diagnosis of primary ciliary dyskinesia and reveals novel candidate genes and genetic variants. PLoS ONE, 2018, 13, e0205422.	2.5	25
20	Variants in TPMT, ITPA, ABCC4 And ABCB1 Genes as Predictors of 6-Mercaptopurine Induced Toxicity in Children with Acute Lymphoblastic Leukemia. Journal of Medical Biochemistry, 2018, 37, 320-327.	1.7	25
21	HLA genotyping in pediatric celiac disease patients. Bosnian Journal of Basic Medical Sciences, 2014, 14, 171-176.	1.0	25
22	Molecular Genetics and Genotype-Based Estimation of BH4-Responsiveness in Serbian PKU Patients: Spotlight on Phenotypic Implications of p.L48S. JIMD Reports, 2012, 9, 49-58.	1.5	22
23	Functional analysis of a novel KLF1 gene promoter variation associated with hereditary persistence of fetal hemoglobin. Annals of Hematology, 2013, 92, 53-58.	1.8	22
24	Optimization of PCR Conditions for Amplification of GC-RichEGFRPromoter Sequence. Journal of Clinical Laboratory Analysis, 2013, 27, 487-493.	2.1	22
25	Expression pattern of long non-coding RNA growth arrest-specific 5 in the remission induction therapy in childhood acute lymphoblastic leukemia. Journal of Medical Biochemistry, 2019, 38, 292-298.	1.7	22
26	Metabolic Syndrome in Inflammatory Bowel Disease: Association with Genetic Markers of Obesity and Inflammation. Metabolic Syndrome and Related Disorders, 2020, 18, 31-38.	1.3	22
27	Functional prediction and comparative population analysis of variants in genes for proteases and innate immunity related to SARS-CoV-2 infection. Infection, Genetics and Evolution, 2020, 84, 104498.	2.3	22
28	Association of Vitamin D, Zinc and Selenium Related Genetic Variants With COVID-19 Disease Severity. Frontiers in Nutrition, 2021, 8, 689419.	3.7	22
29	PARP-1 and YY1 Are Important Novel Regulators of CXCL12 Gene Transcription in Rat Pancreatic Beta Cells. PLoS ONE, 2013, 8, e59679.	2.5	22
30	Polymorphisms of tumor-necrosis factor-Î±â˜308 and lymphotoxin-Î± + 250: Possible modulation of susceptibility to apoptosis in chronic lymphocytic leukemia and non-Hodgkin lymphoma mononuclear cells. Leukemia and Lymphoma, 2008, 49, 2163-2169.	1.3	21
31	6-mercaptopurine influences TPMT gene transcription in a TPMT gene promoter variable number of tandem repeats-dependent manner. Pharmacogenomics, 2012, 13, 283-295.	1.3	21
32	TPMT gene expression is increased during maintenance therapy in childhood acute lymphoblastic leukemia patients in a TPMT gene promoter variable number of tandem repeat-dependent manner. Pharmacogenomics, 2015, 16, 1701-1712.	1.3	21
33	The use of canine mesenchymal stem cells for the autologous treatment of osteoarthritis. Acta Veterinaria Hungarica, 2018, 66, 376-389.	0.5	19
34	Influence Of Promoter Polymorphisms Of The Tnf-Î± (-308g/A) And IL-6 (-174g/C) Genes On Therapeutic Response To Etanercept In Rheumatoid Arthritis. Journal of Medical Biochemistry, 2015, 34, 414-421.	1.7	18
35	Importance of TLR9-IL23-IL17 axis in inflammatory bowel disease development: Gene expression profiling study. Clinical Immunology, 2018, 197, 86-95.	3.2	18
36	Association of gene variants in TLR4 and IL-6 genes with Perthes disease. Srpski Arhiv Za Celokupno Lekarstvo, 2014, 142, 450-456.	0.2	18

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37	Prognostic significance of SOX2, SOX3, SOX11, SOX14 and SOX18 gene expression in adult de novo acute myeloid leukemia. Leukemia Research, 2018, 67, 32-38.	0.8	17
38	Acute myeloid leukemia with NUP98â€“HOXC13 fusion and FLT3 internal tandem duplication mutation: case report and literature review. Cancer Genetics and Cytogenetics, 2009, 193, 98-103.	1.0	16
39	Transcriptional regulation and pharmacogenomics. Pharmacogenomics, 2011, 12, 655-673.	1.3	16
40	Somatic mutations of isocitrate dehydrogenases 1 and 2 are prognostic and follow-up markers in patients with acute myeloid leukaemia with normal karyotype. Radiology and Oncology, 2016, 50, 385-393.	1.7	16
41	The influence of novel transcriptional regulatory element in intron 14 on the expression of Janus kinase 2 gene in myeloproliferative neoplasms. Journal of Applied Genetics, 2013, 54, 21-26.	1.9	15
42	Predictive genetic markers of coagulation, inflammation and apoptosis in Perthes diseaseâ€”Serbian experience. European Journal of Pediatrics, 2015, 174, 1085-1092.	2.7	15
43	Genetic and environmental factors significant for the presentation and development of inflammatory bowel disease. European Journal of Gastroenterology and Hepatology, 2017, 29, 909-915.	1.6	15
44	Pharmacogenomic Markers of Methotrexate Response in the Consolidation Phase of Pediatric Acute Lymphoblastic Leukemia Treatment. Genes, 2020, 11, 468.	2.4	15
45	Distinct Genetic Lesions Drive Leukemogenesis in Secondary Acute Myeloid Leukemia,. Blood, 2011, 118, 3559-3559.	1.4	15
46	Incidence of FLT3 and nucleophosmin gene mutations in childhood acute myeloid leukemia: Serbian experience and the review of the literature. Medical Oncology, 2010, 27, 640-645.	2.5	14
47	Expression of TLR7, TLR9, JAK2, and STAT3 genes in peripheral blood mononuclear cells from patients with systemic sclerosis. Journal of Applied Genetics, 2018, 59, 59-66.	1.9	14
48	Pharmacogenomic markers of glucocorticoid response in the initial phase of remission induction therapy in childhood acute lymphoblastic leukemia. Radiology and Oncology, 2018, 52, 296-306.	1.7	14
49	Variations in inflammatory genes as molecular markers for prediction of inflammatory bowel disease occurrence. Journal of Digestive Diseases, 2015, 16, 723-733.	1.5	13
50	Parallel targeted next generation sequencing of childhood and adult acute myeloid leukemia patients reveals uniform genomic profile of the disease. Tumor Biology, 2016, 37, 13391-13401.	1.8	13
51	Frequencies of EGFR single nucleotide polymorphisms in non-small cell lung cancer patients and healthy individuals in the Republic of Serbia: a preliminary study. Tumor Biology, 2016, 37, 10479-10486.	1.8	13
52	Clinical Applicability of Sequence Variations in Genes Related to Drug Metabolism. Current Drug Metabolism, 2011, 12, 445-454.	1.2	12
53	Prognostic Impact of <i>NPM1</i> Mutations in Serbian Adult Patients with Acute Myeloid Leukemia. Acta Haematologica, 2012, 128, 203-212.	1.4	12
54	Prognostic Significance of Cereblon Expression in Patients With Multiple Myeloma. Clinical Lymphoma, Myeloma and Leukemia, 2016, 16, 610-615.	0.4	12

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55	Impact of alterations in X-linked IRAK1gene and miR-146a on susceptibility and clinical manifestations in patients with systemic sclerosis. Immunology Letters, 2018, 204, 1-8.	2.5	12
56	The importance of combined NGS and MLPA genetic tests for differential diagnosis of maturity onset diabetes of the young. Endokrynologia Polska, 2019, 70, 28-36.	1.0	12
57	Novel transcriptional regulatory element in the phenylalanine hydroxylase gene intron 8. Molecular Genetics and Metabolism, 2010, 101, 81-83.	1.1	11
58	Molecular Genetic Markers as a Basis for Personalized Medicine / MOLEKULARNO-GENETIÄŒKI MARKERI KAO OSNOV ZA PERSONALIZOVANU MEDICINU. Journal of Medical Biochemistry, 2014, 33, 8-21.	1.7	11
59	Genetic predictors of celiac disease, lactose intolerance, and vitamin D function and presence of peptide morphins in urine of children with neurodevelopmental disorders. Nutritional Neuroscience, 2019, 22, 40-50.	3.1	11
60	Clinical relevance of IL-6 gene polymorphism in severely injured patients. Bosnian Journal of Basic Medical Sciences, 2014, 14, 110.	1.0	11
61	Acquired amegakaryocytic thrombocytopenia associated with proliferation of Î³/Î´ TCR+ T-lymphocytes and a BCR-ABL (p210) fusion transcript. European Journal of Haematology, 2004, 73, 372-375.	2.2	10
62	Severe central nervous system thrombotic events in hemoglobin Sabine patient. European Journal of Haematology, 2004, 72, 67-70.	2.2	10
63	Overexpression of the novel member of the BCL2 gene family, BCL2L12, is associated with the disease outcome in patients with acute myeloid leukemia. Clinical Biochemistry, 2012, 45, 1362-1367.	1.9	10
64	Single institute study of FLT3 mutation in acute myeloid leukemia with near tetraploidy in Serbia. Journal of Genetics, 2009, 88, 149-152.	0.7	9
65	Thalassemia Syndromes in Serbia: An update. Hemoglobin, 2010, 34, 477-485.	0.8	9
66	Gammopathy and B lymphocyte clonality in patients with Gaucher type I disease. Blood Cells, Molecules, and Diseases, 2013, 50, 222-225.	1.4	9
67	Novel genetic risk variants for pediatric celiac disease. Human Genomics, 2016, 10, 34.	2.9	9
68	Genes and metabolic pathway of sarcoidosis: identification of key players and risk modifiers. Archives of Medical Science, 2019, 15, 1138-1146.	0.9	9
69	Meta-Analysis of Circulating Cell-Free DNAâ€™s Role in the Prognosis of Pancreatic Cancer. Cancers, 2021, 13, 3378.	3.7	9
70	Machine Learning Modeling from Omics Data as Prospective Tool for Improvement of Inflammatory Bowel Disease Diagnosis and Clinical Classifications. Genes, 2021, 12, 1438.	2.4	9
71	Complex Patterns of Chromosome 11 Aberrations in Myeloid Malignancies Target CBL, MLL, DDB1 and LMO2. PLoS ONE, 2013, 8, e77819.	2.5	9
72	Pharmacotranscriptomic Biomarkers in Glucocorticoid Treatment of Pediatric Inflammatory Bowel Disease. Current Medicinal Chemistry, 2018, 25, 2855-2871.	2.4	9

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73	Relevance of TNF-Î±, IL-6 and IRAK1 gene expression for assessing disease severity and therapy effects in tuberculosis patients. Journal of Infection in Developing Countries, 2019, 13, 419-425.	1.2	9
74	Arterial Ischemic Stroke in a Child with Î²-Thalassemia Trait and Methylentetrahydrofolate Reductase Mutation. Journal of Child Neurology, 2007, 22, 208-210.	1.4	8
75	The Role of Lymphocyte to Monocyte Ratio, Microvessel Density and HiGH CD44 Tumor Cell Expression in Non Hodgkin Lymphomas. Pathology and Oncology Research, 2016, 22, 567-577.	1.9	8
76	Impact of genotype on neutropenia in a large cohort of Serbian patients with glycogen storage disease type Ib. European Journal of Medical Genetics, 2020, 63, 103767.	1.3	8
77	Patients with early stage chronic lymphocytic leukemia: new risk stratification based on molecular profiling. Leukemia and Lymphoma, 2011, 52, 1394-1397.	1.3	7
78	The predictive value of morphological findings in early diagnosis of acute myeloid leukemia with recurrent cytogenetic abnormalities. Leukemia Research, 2018, 75, 23-28.	0.8	7
79	The influence of Wilms' tumor 1 gene expression level on prognosis and risk stratification of acute promyelocytic leukemia patients. International Journal of Laboratory Hematology, 2020, 42, 82-87.	1.3	7
80	The FKBP5 genotype and childhood trauma effects on FKBP5 DNA methylation in patients with psychosis, their unaffected siblings, and healthy controls. Psychoneuroendocrinology, 2021, 128, 105205.	2.7	7
81	IL-28B genotypes as predictors of long-term outcome in patients with hepatitis C-related severe liver injury. Journal of Infection in Developing Countries, 2019, 13, 526-535.	1.2	7
82	Association of gene variants in TLR4 and IL-6 genes with Perthes disease. Srpski Arhiv Za Celokupno Lekarstvo, 2014, 142, 450-6.	0.2	7
83	Compound heterozygosity for the Cretan type of non-deletional hereditary persistence of fetal hemoglobin and Î²-thalassemia or Hb Sabine confirms the functional role of the AÎ³ â^'158 C>T mutation in Î³-globin gene transcription. Blood Cells, Molecules, and Diseases, 2008, 41, 263-264.	1.4	6
84	Effects of DMSO, glycerol, betaine and their combinations in detecting single nucleotide polymorphisms of epidermal growth factor receptor (EGFR) gene promoter sequence in non-small-cell lung cancer (NSCLC) patients. Journal of Pharmaceutical and Biomedical Analysis, 2016, 128, 275-279.	2.8	6
85	Expression Pattern and Prognostic Significance of EVI1 Gene in Adult Acute Myeloid Leukemia Patients with Normal Karyotype. Indian Journal of Hematology and Blood Transfusion, 2020, 36, 292-299.	0.6	6
86	Pharmacogenomics landscape of COVID-19 therapy response in Serbian population and comparison with worldwide populations. Journal of Medical Biochemistry, 2020, 39, 488-499.	1.7	6
87	Clinical Application of Thiopurine Pharmacogenomics in Pediatrics. Current Drug Metabolism, 2020, 21, 53-62.	1.2	6
88	FLT3/D835 mutation and inversion of chromosome 16 in leukemic transformation of myelofibrosis. European Journal of Internal Medicine, 2006, 17, 434-435.	2.2	5
89	Novel Patched 1 mutations in patients with nevoid basal cell carcinoma syndrome â€“ case report. Croatian Medical Journal, 2015, 56, 63-67.	0.7	5
90	Functional Analysis of an^AÎ³-Globin Gene Promoter Variant (HBG1) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 67 Td (g.-2 Stress. Hemoglobin, 2016, 40, 48-52.	0.8	5

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91	CRISPR/Cas9 genome editing of SLC37A4 gene elucidates the role of molecular markers of endoplasmic reticulum stress and apoptosis in renal involvement in glycogen storage disease type Ib. Gene, 2019, 703, 17-25.	2.2	5
92	Molecular Characteristics, Phenotypic Diversity and Genotype-Estimated Therapeutic Responsiveness of Serbian Patients with Phenylketonuria / MOLEKULARNE KARAKTERISTIKE, FENOTIPSKA RAZNOLIKOST I PROCENA ODGOVORA NA TERAPIJU ZASNOVANA NA GENOTIPU KOD SRPSKIH PACIJENATA SA FENILKETONURIJOM. Journal of Medical Biochemistry, 2014, 33, 97-107.	1.7	4
93	Association between the -174 C/G polymorphism in the interleukin-6 (IL-6) gene and gastrointestinal involvement in patients with systemic sclerosis. Clinical Rheumatology, 2018, 37, 2447-2454.	2.2	4
94	Complex transcriptional regulation of the BCL2L12 gene: Novel, active promoter in K562 cells. Gene, 2020, 750, 144723.	2.2	4
95	Identification and Classification of Novel Genetic Variants: En Route to the Diagnosis of Primary Ciliary Dyskinesia. International Journal of Molecular Sciences, 2021, 22, 8821.	4.1	4
96	Mutations in the PAH gene: A Tool for population genetics study. Archives of Biological Sciences, 2007, 59, 161-167.	0.5	4
97	Expression Profiles of Long Non-Coding RNA GAS5 and MicroRNA-222 in Younger AML Patients. Diagnostics, 2022, 12, 86.	2.6	4
98	Expression pattern of circulating long non-coding RNA GAS5 as a novel biomarker in non-small cell lung cancer patients. Archives of Medical Science, 2020, , .	0.9	3
99	Exploring inflammatory and apoptotic signatures in distinct Crohn's disease phenotypes: Way towards molecular stratification of patients and targeted therapy. Pathology Research and Practice, 2020, 216, 152945.	2.3	3
100	Genetic variants in TNFA, LTA, TLR2 and TLR4 genes and risk of sepsis in patients with severe trauma: nested case-control study in a level-1 trauma centre in SERBIA. Injury, 2021, 52, 419-425.	1.7	3
101	Immunoglobulin Heavy Chain Gene Rearrangements in Patients with Gaucher Disease. Journal of Medical Biochemistry, 2018, 37, 307-312.	1.7	3
102	Significance of UGT1A1*28 Genotype in Patients with Advanced Liver Injury Caused by Chronic Hepatitis C. Journal of Medical Biochemistry, 2019, 38, 45-52.	1.7	3
103	Functional Characterization of Novel Phenylalanine Hydroxylase p.Gln226Lys Mutation Revealed Its Non-responsiveness to Tetrahydrobiopterin Treatment in Hepatoma Cellular Model. Biochemical Genetics, 2018, 56, 533-541.	1.7	2
104	Association of SLC28A3 Gene Expression and CYP2B6*6 Allele with the Response to Fludarabine Plus Cyclophosphamide in Chronic Lymphocytic Leukemia Patients. Pathology and Oncology Research, 2020, 26, 743-752.	1.9	2
105	Analysis of the promoter regions of disease-causing genes in maturity-onset diabetes of the young patients. Molecular Biology Reports, 2020, 47, 6759-6768.	2.3	2
106	Next-Generation Sequencing: The Enabler and the Way Ahead. , 2020, , 175-200.		2
107	Case Report: Successful Therapy of Spontaneously Occurring Canine Degenerative Lumbosacral Stenosis Using Autologous Adipose Tissue-Derived Mesenchymal Stem Cells. Frontiers in Veterinary Science, 2021, 8, 732073.	2.2	2
108	Î±-Tocopherol Acetate Attenuates Mitochondrial Oxygen Consumption and Maintains Primitive Cells within Mesenchymal Stromal Cell Population. Stem Cell Reviews and Reports, 2021, 17, 1390-1405.	3.8	2

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109	Next generation sequencing as a tool for pharmacogenomic profiling: Nine novel potential genetic markers for targeted therapy in childhood acute lymphoblastic leukemia. Srpski Arhiv Za Celokupno Lekarstvo, 2018, 146, 407-411.	0.2	2
110	Genomic profiling of thymoma using a targeted high-throughput approach. Archives of Medical Science, 2020, , .	0.9	2
111	Association of Mitochondrial DNA Variants and Cognitive Impairment of Phenylketonuria Patients / POVEZANOST VARIJANTI U MITOHONDRIJALNOJ DNK I KOGNITIVNOG FENOTIPA KOD PACIJENATA SA FENILKETONURIJOM. Journal of Medical Biochemistry, 2013, 32, 347-353.	1.7	1
112	Expression Of The Bcl2 Gene In Chronic Lymphocytic Leukaemia Patients. Serbian Journal of Experimental and Clinical Research, 2015, 16, 187-191.	0.1	1
113	New PAH gene promoter KLF1 and 3â€²-region C/EBPalpha motifs influence transcription in vitro. Journal of Applied Genetics, 2017, 58, 79-85.	1.9	1
114	Prognostic significance of combined BAALC and MN1 gene expression level in acute myeloid leukemia with normal karyotype. International Journal of Laboratory Hematology, 2021, 43, 433-440.	1.3	1
115	The first insight into the genetic structure of the population of modern Serbia. Scientific Reports, 2021, 11, 13995.	3.3	1
116	Intra-articular injection of autologous adipose-derived mesenchymal stem cells in the treatment of knee osteoarthritis. , 2018, 20, e3002.		1
117	Reappraising Immunoglobulin Repertoire Restrictions in Chronic Lymphocytic Leukemia: Focus on Major Stereotyped Subsets and Closely Related Satellites. Blood, 2016, 128, 4376-4376.	1.4	1
118	Biochemical phenotype and origin of the three most common beta-thalassemia mutations in Serbia. Journal of Medical Biochemistry, 2004, 23, 361-366.	0.1	1
119	TPMT gene polymorphisms: on the doorstep of personalized medicine. Indian Journal of Medical Research, 2009, 129, 478-80.	1.0	1
120	Novel Therapy Approaches in Î²-Thalassemia Syndromes â€" A Role of Genetic Modifiers. , 2015, , .		0
121	PERSONALIZATION OF 6-MERCAPTOPURINE THERAPY FOR CHILDREN WITH ACUTE LYMPHOBLASTIC LEUKEMIA. Archives of Disease in Childhood, 2016, 101, e1.5-e1.	1.9	0
122	A novel 9â€‰bp deletion (c.1271_1279delGTGCCCGCG) in exon 10 of CYP21A2 gene causing severe congenital adrenal hyperplasia. Endocrine, 2021, 73, 196-202.	2.3	0
123	Application of targeted next generation sequencing for the mutational profiling of patients with acute lymphoblastic leukemia. Journal of Medical Biochemistry, 2019, 39, 72-82.	1.7	0
124	The pharmacogenomics of vincristine-induced peripheral neuropathy in pediatric acute lymphoblastic leukemia patients in Serbia - a single center experience. Srpski Arhiv Za Celokupno Lekarstvo, 2022, 150, 53-58.	0.2	0
125	High risk population screening for Fabry disease in hemodialysis patients in Vojvodina: Pilot study. Srpski Arhiv Za Celokupno Lekarstvo, 2022, 150, 281-287.	0.2	0