Evgeny N Suspitsin

List of Publications by Year
in descending order

60 papers	1,128 citations	430754 18 h-index	414303 32 g-index
60 all docs	60 docs citations	60 times ranked	1947 citing authors

#	Article	IF	Citations
1	Bardet-Biedl Syndrome. Molecular Syndromology, 2016, 7, 62-71.	0.3	103
2	High response rates to neoadjuvant platinum-based therapy in ovarian cancer patients carrying germ-line BRCA mutation. Cancer Letters, 2015, 369, 363-367.	3.2	82
3	High frequency of BRCA1 5382insC mutation in Russian breast cancer patients. European Journal of Cancer, 2006, 42, 1380-1384.	1.3	70
4	Evidence against involvement of p53 polymorphism in breast cancer predisposition. International Journal of Cancer, 2003, 103, 431-433.	2.3	65
5	Concordance of allelic imbalance profiles in synchronous and metachronous bilateral breast carcinomas. International Journal of Cancer, 2002, 100, 557-564.	2.3	64
6	NBS1 657del5 mutation may contribute only to a limited fraction of breast cancer cases in Russia. International Journal of Cancer, 2005, 114, 585-589.	2.3	59
7	High prevalence and breast cancer predisposing role of the BLM c.1642 C>T (Q548X) mutation in Russia. International Journal of Cancer, 2012, 130, 2867-2873.	2.3	58
8	CYP19 gene polymorphism in endometrial cancer patients. Journal of Cancer Research and Clinical Oncology, 2001, 127, 135-138.	1.2	51
9	Pattern of clinically relevant mutations in consecutive series of Russian colorectal cancer patients. Medical Oncology, 2013, 30, 686.	1.2	43
10	Multigene sequencing reveals heterogeneity of NLRP12-related autoinflammatory disorders. Rheumatology International, 2018, 38, 887-893.	1.5	42
11	Non-founder BRCA1 mutations in Russian breast cancer patients. Cancer Letters, 2010, 298, 258-263.	3.2	34
12	High frequency of BRCA1, but not CHEK2 or NBS1 (NBN), founder mutations in Russian ovarian cancer patients. Hereditary Cancer in Clinical Practice, 2009, 7, 5.	0.6	31
13	High prevalence ofGPRC5Agermline mutations inBRCA1-mutant breast cancer patients. International Journal of Cancer, 2014, 134, 2352-2358.	2.3	31
14	Identification of novel hereditary cancer genes by whole exome sequencing. Cancer Letters, 2015, 369, 274-288.	3.2	31
15	CYP17 polymorphism in the groups of distinct breast cancer susceptibility: comparison of patients with the bilateral disease vs. monolateral breast cancer patients vs. middle-aged female controls vs. elderly tumor-free women. Cancer Letters, 2000, 156, 45-50.	3.2	28
16	Gene rearrangements in consecutive series of pediatric inflammatory myofibroblastic tumors. Pediatric Blood and Cancer, 2020, 67, e28220.	0.8	24
17	Evidence for microsatellite instability in bilateral breast carcinomas. Cancer Letters, 2000, 154, 9-17.	3.2	20
18	Development of breast tumors in CHEK2, NBN/NBS1 and BLM mutation carriers does not commonly involve somatic inactivation of the wild-type allele. Medical Oncology, 2014, 31, 828.	1.2	20

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19	Distinct prevalence of the CYP19 Î”3(TTTA)7 allele in premenopausal versus postmenopausal breast cancer patients, but not in control individuals. European Journal of Cancer, 2002, 38, 1911-1916.	1.3	19
20	Microsatellite instability analysis of bilateral breast tumors suggests treatment-related origin of some contralateral malignancies. Journal of Cancer Research and Clinical Oncology, 2006, 133, 57-64.	1.2	18
21	High sensitivity of BRCA1-associated tumors to cisplatin monotherapy: report of two cases. Cancer Genetics and Cytogenetics, 2010, 197, 91-94.	1.0	18
22	Mixed epithelial/mesenchymal metaplastic carcinoma (carcinosarcoma) of the breast in BRCA1 carrier. Breast Cancer, 2011, 18, 137-140.	1.3	18
23	Wholeâ€‘exome sequencing in Russian children with nonâ€‘type 1 diabetes mellitus reveals a wide spectrum of genetic variants in MODYâ€‘related and unrelated genes. Molecular Medicine Reports, 2019, 20, 4905-4914.	1.1	18
24	Detection of BRCA1 gross rearrangements by droplet digital PCR. Breast Cancer Research and Treatment, 2017, 165, 765-770.	1.1	16
25	Spectrum of APC and MUTYH germâ€line mutations in Russian patients with colorectal malignancies. Clinical Genetics, 2018, 93, 1015-1021.	1.0	16
26	Paired distribution of molecular subtypes in bilateral breast carcinomas. Cancer Genetics, 2011, 204, 96-102.	0.2	14
27	ATM mutation spectrum in Russian children with ataxia-telangiectasia. European Journal of Medical Genetics, 2020, 63, 103630.	0.7	13
28	Nonrandom distribution of oncogene amplifications in bilateral breast carcinomas: Possible role of host factors and survival bias. International Journal of Cancer, 2007, 120, 297-302.	2.3	10
29	Next generation sequencing analysis of consecutive Russian patients with clinical suspicion of inborn errors of immunity. Clinical Genetics, 2020, 98, 231-239.	1.0	10
30	The spectrum of Lynch syndrome-associated germ-line mutations in Russia. European Journal of Medical Genetics, 2020, 63, 103753.	0.7	9
31	Exome sequencing study of Russian breast cancer patients suggests a predisposing role for USP39. Breast Cancer Research and Treatment, 2020, 179, 731-742.	1.1	9
32	Exome Sequencing of a Family with Bardet-Biedl Syndrome Identifies the Common Russian Mutation c.1967_1968delTAinsC in <i>BBS7</i>. Molecular Syndromology, 2015, 6, 96-98.	0.3	8
33	L-myc polymorphism in cancer patients, healthy blood donors and elderly, tumor-free individuals in Russia. , 2000, 85, 747-750.		7
34	Apoptosis-deficient Pro allele of gene is associated with the resistance of psoriasis to the UV-based therapy. Journal of Dermatological Science, 2005, 37, 185-187.	1.0	7
35	CHEK2 1100 delC mutation in Russian ovarian cancer patients. Hereditary Cancer in Clinical Practice, 2007, 5, 153.	0.6	7
36	Pattern of TSC1 and TSC2 germline mutations in Russian patients with tuberous sclerosis. Journal of Human Genetics, 2018, 63, 597-604.	1.1	7

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37	Exome-based search for recurrent disease-causing alleles in Russian population. European Journal of Medical Genetics, 2019, 62, 103656.	0.7	7
38	TP53 mutations in synchronous and metachronous bilateral breast carcinomas. Cancer Genetics and Cytogenetics, 2008, 184, 119-121.	1.0	6
39	First Two Cases of Bloom Syndrome in Russia: Lack of Skin Manifestations in a BLM c.1642C>T (p.Q548X) Homozygote as a Likely Cause of Underdiagnosis. Molecular Syndromology, 2017, 8, 103-106.	0.3	6
40	Evidence for angiogenesis-independent contribution of VEGFR1 (FLT1) in gastric cancer recurrence. Medical Oncology, 2013, 30, 644.	1.2	4
41	PREDICTIVE RESPONSE MARKERS FOR IMMUNE RESPONSE BLOCKS. Siberian Journal of Oncology, 2020, 19, 123-131.	0.1	4
42	Improved Reliability of Allele-Specific PCR. BioTechniques, 2002, 33, 484-490.	0.8	4
43	Large family with both parents affected by distinct BRCA1 mutations: implications for genetic testing. Hereditary Cancer in Clinical Practice, 2009, 7, 2.	0.6	3
44	Rapid Symptomatic Improvement in Gefitinib-Treated Patients with EGFR-Mutated Lung Cancer: Possible Role of Downregulation of Inflammatory Molecules?. Onkologie, 2011, 34, 559-560.	1.1	3
45	Systemic Lupus Erythematosus with neuropsychiatric manifestations in a child: case report and review of the international recommendations for the diagnostics and management. Rossiyskiy Vestnik Perinatologii I Pediatrii, 2021, 66, 98-105.	0.1	3
46	Diagnosis of carcinoma of unknown primary site with the aid of simple PCR tests: a single-center experience. Neoplasma, 2018, 65, 461-468.	0.7	2
47	Evidence for a pathogenic role of BRCA1 L1705P and W1837X germ-line mutations. Molecular Biology Reports, 2016, 43, 335-338.	1.0	1
48	Revisiting multiple erroneous genetic testing results and clinical misinterpretations in a patient with Li-Fraumeni syndrome: lessons for translational medicine. Hereditary Cancer in Clinical Practice, 2021, 19, 2.	0.6	1
49	HEREDITARY ANGIOEDEMA â€“ A PROBLEM AT THE INTERSECTION OF IMMUNOLOGY AND ALLERGOLOGY: ANALYSIS OF LITERATURE DATA AND DESCRIPTION OF A SERIES OF 34 CASES. Pediatriia, 2021, 100, 49-57.	0.1	1
50	MEVALONATE KINASE DEFICIENCY IN AN INFANT: KEY ASPECTS OF DIAGNOSIS AND TREATMENT. Pediatriia, 2021, 100, 276-283.	0.1	1
51	Analysis of interferon type I signature for differential diagnosis of diseases of the immune system () Tj ETQq1 1 0.784314 rgBT /Overlo	0.2	1
52	Primary tumor (tumoral) calcification is a rare disease in the practice of a rheumatologist and orthopedist: experience with the use of an interleukin-1 inhibitor in combination with surgical correction. Pediatric Traumatology, Orthopaedics and Reconstructive Surgery, 2019, 7, 85-92.	0.1	1
53	GP134â€...Confirmation of pathogenetic heterogeneity of diabetes mellitus in children using whole-exome sequencing. , 2019, , .		0
54	ACTIVATED PHOSPHOINOSITIDE 3-KINASE Î´ SYNDROME â€“ A RARE FORM OF PRIMARY IMMUNODEFICIENCY. Pediatriia, 2021, 100, 248-253.	0.1	0

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55	ANALYSIS OF INTERFERON-I SIGNALING ACTIVITY IN CHILDREN WITH SYSTEMIC LUPUS ERYTHEMATOSUS: RESULTS OF A PILOT PROSPECTIVE STUDY. Pediatriia, 2021, 100, 77-88.	0.1	0
56	Abstract 2597: Identification of new genes for hereditary breast cancer. , 2012, , .		0
57	Case report of the inflammatory myofibroblastic tumor of the liver in infant. Pediatric Hematology/Oncology and Immunopathology, 2018, 17, 93-102.	0.1	0
58	RECURRENT STREPTOCOCCAL INFECTION IN A CHILD WITH A CONGENITAL DEFICIENCY OF MANNOSE-BINDING LECTIN. Pediatriia, 2020, 99, 266-270.	0.1	0
59	Identification of recurrent pathogenic alleles using exome sequencing data: Proof-of-concept study of Russian subjects. European Journal of Medical Genetics, 2022, 65, 104426.	0.7	0
60	Genetic variants identified in children with recurrent infections. Detskie Infekcii (Moskva), 2021, 20, 13-17.	0.1	0