Grzegorz Kurzawski

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	A genetic variant in telomerase reverse transcriptase (TERT) modifies cancer risk in Lynch syndrome patients harbouring pathogenic MSH2 variants. Scientific Reports, 2021, 11, 11401.	3.3	6
2	CD36 gene polymorphism and plasma sCD36 as the risk factor in higher cholesterolemia. Archives De Pediatrie, 2018, 25, 177-181.	1.0	10
3	New <i><scp>EPCAM</scp></i> founder deletion in Polish population. Clinical Genetics, 2017, 92, 649-653.	2.0	6
4	Cumulative Small Effect Genetic Markers and the Risk of Colorectal Cancer in Poland, Estonia, Lithuania, and Latvia. Gastroenterology Research and Practice, 2015, 2015, 1-10.	1.5	5
5	Management of ovarian and endometrial cancers in women belonging to HNPCC carrier families: review of the literature and results of cancer risk assessment in Polish HNPCC families. Hereditary Cancer in Clinical Practice, 2015, 13, 3.	1.5	11
6	Is plasma soluble CD36 associated with cardiovascular risk factors in early onset coronary artery disease patients?. Scandinavian Journal of Clinical and Laboratory Investigation, 2015, 75, 398-406.	1.2	18
7	Polymorphisms in nucleotide excision repair genes and susceptibility to colorectal cancer in the Polish population. Molecular Biology Reports, 2015, 42, 755-764.	2.3	32
8	Cumulative effects of genetic markers and the detection of advanced colorectal neoplasias by population screening. Clinical Genetics, 2015, 88, 234-240.	2.0	5
9	Lynch syndrome mutations shared by the Baltic States and Poland. Clinical Genetics, 2014, 86, 190-193.	2.0	5
10	Germline deletions in the EPCAM gene as a cause of Lynch syndrome – literature review. Hereditary Cancer in Clinical Practice, 2013, 11, 9.	1.5	104
11	Combined analysis of three lynch syndrome cohorts confirms the modifying effects of 8q23.3 and 11q23.1 in MLH1 mutation carriers. International Journal of Cancer, 2013, 132, 1556-1564.	5.1	33
12	ls CD36 gene polymorphism in region encoding lipid-binding domain associated with early onset CAD?. Gene, 2013, 530, 134-137.	2.2	6
13	Association of CD36 gene polymorphisms with echo- and electrocardiographic parameters in patients with early onset coronary artery disease. Archives of Medical Science, 2013, 4, 640-650.	0.9	7
14	DNA repair gene polymorphisms and risk of early onset colorectal cancer in Lynch syndrome. Cancer Epidemiology, 2012, 36, 183-189.	1.9	25
15	DNA and RNA analyses in detection of genetic predisposition to cancer. Hereditary Cancer in Clinical Practice, 2012, 10, 17.	1.5	6
16	Fast diagnostic test for the identification of an increased genetic predisposition to colon cancer (exemplified on a DNA test for recurrent mutations of the gene MMR). Hereditary Cancer in Clinical Practice, 2012, 10, A13.	1.5	0
17	Polymorphism of the CD36 Gene and Cardiovascular Risk Factors in Patients with Coronary Artery Disease Manifested at a Young Age. Biochemical Genetics, 2012, 50, 103-111.	1.7	22
18	Polymorphism of CD36 gene, carbohydrate metabolism and plasma CD36 concentration in obese children. A preliminary study. Postepy Higieny I Medycyny Doswiadczalnej, 2012, 66, 954-958.	0.1	5

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19	CD36 gene is associated with thickness of atheromatous plaque and ankle-brachial index in patients with early coronary artery disease. Kardiologia Polska, 2012, 70, 918-23.	0.6	12
20	Association of MMP8 gene variation with an increased risk of malignant melanoma. Melanoma Research, 2011, 21, 464-468.	1.2	19
21	AMPD1 gene mutations are associated with obesity and diabetes in Polish patients with cardiovascular diseases. Journal of Applied Genetics, 2011, 52, 67-76.	1.9	16
22	Colorectal cancer susceptibility loci on chromosome 8q23.3 and 11q23.1 as modifiers for disease expression in lynch syndrome. Journal of Medical Genetics, 2011, 48, 279-284.	3.2	44
23	BRCA1 mutations and colorectal cancer in Poland. Familial Cancer, 2010, 9, 541-544.	1.9	33
24	Variant alleles of the CYP1B1 gene are associated with colorectal cancer susceptibility. BMC Cancer, 2010, 10, 420.	2.6	28
25	CHEK2 mutations and HNPCCâ€related colorectal cancer. International Journal of Cancer, 2010, 126, 3005-3009.	5.1	28
26	Analysis of Human <i>CD36</i> Gene Sequence Alterations in the Oxidized Low-Density Lipoprotein-Binding Region Using Denaturing High-Performance Liquid Chromatography. Genetic Testing and Molecular Biomarkers, 2010, 14, 551-557.	0.7	6
27	Combined iPLEX and TaqMan Assays to Screen for 45 Common Mutations in Lynch Syndrome and FAP Patients. Journal of Molecular Diagnostics, 2010, 12, 82-90.	2.8	8
28	Nonalcoholic fatty liver disease and <i>HFE</i> gene mutations: A Polish study. World Journal of Gastroenterology, 2010, 16, 2531.	3.3	29
29	HFE gene mutations in patients with alcoholic liver disease. A prospective study from northwestern Poland. , 2010, 120, 127-31.		2
30	Haemochromatosis <i>HFE</i> gene polymorphisms as potential modifiers of hereditary nonpolyposis colorectal cancer risk and onset age. International Journal of Cancer, 2009, 125, 78-83.	5.1	39
31	MTHFR 677 C>T and 1298 A>C polymorphisms and the age of onset of colorectal cancer in hereditary nonpolyposis colorectal cancer. European Journal of Human Genetics, 2009, 17, 629-635.	2.8	17
32	Clinical characteristics of tumors derived from colorectal cancer patients who harbor the Tumor Necrosis Factor α-1031T/T and NOD2 3020insC polymorphism. Cancer Epidemiology, 2009, 33, 161-163.	1.9	2
33	Frequency of mutations related to hereditary haemochromatosis in northwestern Poland. Journal of Applied Genetics, 2008, 49, 105-107.	1.9	10
34	Auroraâ€A and Cyclin D1 polymorphisms and the age of onset of colorectal cancer in hereditary nonpolyposis colorectal cancer. International Journal of Cancer, 2008, 122, 1273-1277.	5.1	28
35	IGF1 is a modifier of disease risk in hereditary nonâ€polyposis colorectal cancer. International Journal of Cancer, 2008, 123, 1339-1343.	5.1	25
36	Inflammatory response gene polymorphisms and their relationship with colorectal cancer risk. BMC Cancer, 2008, 8, 112.	2.6	41

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37	The â^'149C>T SNP within the ΔDNMT3B gene, is not associated with early disease onset in hereditary non-polyposis colorectal cancer. Cancer Letters, 2008, 265, 39-44.	7.2	10
38	DNA and RNA analyses in detection of genetic predisposition to cancer. Hereditary Cancer in Clinical Practice, 2008, 6, 73.	1.5	0
39	MSH2 and MLH1 testing. Hereditary Cancer in Clinical Practice, 2008, 6, 83.	1.5	Ο
40	DNA testing for variants conferring low or moderate increase in the risk of cancer. Hereditary Cancer in Clinical Practice, 2008, 6, 84.	1.5	3
41	CARD15 variants in patients with sporadic Parkinson's disease. Neuroscience Research, 2007, 57, 473-476.	1.9	67
42	Germline CHEK2 mutations and colorectal cancer risk: different effects of a missense and truncating mutations?. European Journal of Human Genetics, 2007, 15, 237-241.	2.8	61
43	Nationwide study of clinical and molecular features of hereditary non-polyposis colorectal cancer (HNPCC) in Latvia. Anticancer Research, 2007, 27, 653-8.	1.1	4
44	Low-risk Genes and Multi-organ Cancer Risk in the Polish Population. Hereditary Cancer in Clinical Practice, 2006, 4, 52.	1.5	1
45	Some aspects of molecular diagnostics in Lynch syndrome. Hereditary Cancer in Clinical Practice, 2006, 4, 197.	1.5	1
46	Frequency and nature of hMSH6 germline mutations in Polish patients with colorectal, endometrial and ovarian cancers. Clinical Genetics, 2006, 70, 68-70.	2.0	6
47	Prevalence of the NOD2 3020insC mutation in aggregations of breast and lung cancer. Breast Cancer Research and Treatment, 2006, 95, 141-145.	2.5	31
48	CDKN2A common variant and multi-organ cancer risk—a population-based study. International Journal of Cancer, 2006, 118, 3180-3182.	5.1	26
49	MC1R common variants, CDKN2A and their association with melanoma and breast cancer risk. International Journal of Cancer, 2006, 119, 2597-2602.	5.1	38
50	Genetic Polymorphisms in Xenobiotic Clearance Genes and Their Influence on Disease Expression in Hereditary Nonpolyposis Colorectal Cancer Patients. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 2307-2310.	2.5	23
51	The 3020insC Allele of NOD2 Predisposes to Cancers of Multiple Organs. Hereditary Cancer in Clinical Practice, 2005, 3, 59.	1.5	26
52	NOD2 variants and the risk of malignant melanoma. European Journal of Cancer Prevention, 2005, 14, 143-146.	1.3	16
53	Germline MSH2 and MLH1 mutational spectrum including large rearrangements in HNPCC families from Poland (update study). Clinical Genetics, 2005, 69, 40-47.	2.0	34
54	Polymorphism in the P-glycoprotein drug transporter MDR1 gene in colon cancer patients. European Journal of Clinical Pharmacology, 2005, 61, 389-394.	1.9	79

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55	The 3020insC allele of NOD2 predisposes to early-onset breast cancer. Breast Cancer Research and Treatment, 2005, 89, 91-93.	2.5	47
56	CDKN2A common variants and their association with melanoma risk: a population-based study. Cancer Research, 2005, 65, 835-9.	0.9	43
57	Importance of microsatellite instability (MSI) in colorectal cancer: MSI as a diagnostic tool. Annals of Oncology, 2004, 15, iv283-iv284.	1.2	28
58	A Novel Founder CHEK2 Mutation is Associated with Increased Prostate Cancer Risk: Table 1. Cancer Research, 2004, 64, 2677-2679.	0.9	137
59	The NOD2 3020insC Mutation and the Risk of Colorectal Cancer: Table 1. Cancer Research, 2004, 64, 1604-1606.	0.9	105
60	Germline mutation and large deletion analysis of theCDKN2A andARF genes in families with multiple melanoma or an aggregation of malignant melanoma and breast cancer. International Journal of Cancer, 2004, 110, 558-562.	5.1	24
61	CHEK2 Is a Multiorgan Cancer Susceptibility Gene. American Journal of Human Genetics, 2004, 75, 1131-1135.	6.2	426
62	Rarity of germline 1100delC mutation in CHK2 in patients with malignant melanoma of the skin. Melanoma Research, 2004, 14, 121-124.	1.2	8
63	Germline 657del5 mutation in the NBS1 gene in breast cancer patients. International Journal of Cancer, 2003, 106, 379-381.	5.1	80
64	Increased risk of breast cancer in relatives of malignant melanoma patients from families with strong cancer familial aggregation. European Journal of Cancer Prevention, 2003, 12, 241-245.	1.3	14
65	Germline 657del5 mutation in the NBS1 gene in patients with malignant melanoma of the skin. Melanoma Research, 2003, 13, 365-370.	1.2	30
66	Nuclear Pedigree Criteria of Suspected HNPCC. Hereditary Cancer in Clinical Practice, 2003, 1, 1.	1.5	8
67	Electro-Oculographic and Electroretinographic Studies in HNPCC Gene Mutation Carriers. Ophthalmic Research, 2003, 35, 281-294.	1.9	2
68	Germline MSH2 and MLH1 mutational spectrum in HNPCC families from Poland and the Baltic States. Journal of Medical Genetics, 2002, 39, 65e-65.	3.2	26
69	Germline mutations in the von Hippel-Lindau (VHL) gene in patients from Poland: disease presentation in patients with deletions of the entire VHL gene. Journal of Medical Genetics, 2002, 39, 38e-38.	3.2	47
70	Mutation analysis of MLH1 and MSH2 genes performed by denaturing high-performance liquid chromatography. Journal of Proteomics, 2002, 51, 89-100.	2.4	44
71	Ovarian cancer of endometrioid type as part of the MSH6 gene mutation phenotype. Journal of Human Genetics, 2002, 47, 0529-0531.	2.3	12
72	Molecular basis of inherited predispositions for tumors Acta Biochimica Polonica, 2002, 49, 571-581.	0.5	3

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73	Comparison of Alu-PCR, microsatellite instability, and immunohistochemical analyses in finding features characteristic for hereditary nonpolyposis colorectal cancer. Journal of Cancer Research and Clinical Oncology, 2001, 127, 565-569.	2.5	8
74	Fluorescence In Situ Detection of Human Cutaneous Melanoma: Study of Diagnostic Parameters of the Method. Journal of Investigative Dermatology, 2001, 117, 1449-1451.	0.7	26
75	Optimization of experimental conditions for RNA-based sequencing of MLH1 and MSH2 genes. Human Mutation, 2001, 17, 52-60.	2.5	19
76	Value of pedigree/clinical data, immunohistochemistry and microsatellite instability analyses in reducing the cost of determining hMLH1 and hMSH2 gene mutations in patients with colorectal cancer. European Journal of Cancer, 2000, 36, 49-54.	2.8	99
77	Molecular Analyses in Diagnosis of High Genetic Predispositions to Malignancies Journal of Clinical Biochemistry and Nutrition, 2000, 28, 159-165.	1.4	0
78	Relationship between acetylation polymorphism and risk of atopic diseases. Clinical Pharmacology and Therapeutics, 1999, 65, 562-569.	4.7	29
79	Polymorphism of GSTM1 gene in patients with colorectal cancer and colonic polyps. Experimental and Toxicologic Pathology, 1999, 51, 321-325.	2.1	22
80	Frequency and nature of germline Rb-1 gene mutations in a series of patients with sporadic unilateral retinoblastoma. European Journal of Cancer, 1999, 35, 1824-1827.	2.8	12
81	Long polymerase chain reaction in detection of germline deletions in the von Hippel-Lindau tumour suppressor gene. Human Genetics, 1999, 105, 333-336.	3.8	18
82	Age at diagnosis to discriminate those patients for whom constitutional DNA sequencing is appropriate in sporadic unilateral retinoblastoma. European Journal of Cancer, 1998, 34, 1919-1921.	2.8	17
83	Leptin receptor isoforms expressed in human adipose tissue. Metabolism: Clinical and Experimental, 1998, 47, 844-847.	3.4	110
84	Hereditary breast cancer. Polish Journal of Pathology, 1998, 49, 59-66.	0.3	2
85	Losses at 3p common deletion sites in subtypes of kidney tumours: histopathological correlations. Virchows Archiv Fur Pathologische Anatomie Und Physiologie Und Fur Klinische Medizin, 1996, 429, 37-42.	2.8	24
86	Mechanism of the Inhibitory Effect of Curdlan Sulfate on HIV-1 Infection in Vitro. Virology, 1994, 202, 735-745.	2.4	101
87	Analysis of ??+ T cells in peripheral blood of children with perinatal human immunodeficiency virus (HIV) infection, Journal of Clinical Immunology, 1993, 13, 193-203.	3.8	20