Grzegorz Kurzawski

List of Publications by Year
in descending order

87	2,710	185998 28	189595 50
papers	citations	h-index	g-index

88	88	88	3818
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	CHEK2 Is a Multiorgan Cancer Susceptibility Gene. American Journal of Human Genetics, 2004, 75, 1131-1135.	2.6	426
2	A Novel Founder CHEK2 Mutation is Associated with Increased Prostate Cancer Risk: Table 1. Cancer Research, 2004, 64, 2677-2679.	0.4	137
3	Leptin receptor isoforms expressed in human adipose tissue. Metabolism: Clinical and Experimental, 1998, 47, 844-847.	1.5	110
4	The NOD2 3020insC Mutation and the Risk of Colorectal Cancer: Table 1. Cancer Research, 2004, 64, 1604-1606.	0.4	105
5	Germline deletions in the EPCAM gene as a cause of Lynch syndrome â€“ literature review. Hereditary Cancer in Clinical Practice, 2013, 11, 9.	0.6	104
6	Mechanism of the Inhibitory Effect of Curdlan Sulfate on HIV-1 Infection in Vitro. Virology, 1994, 202, 735-745.	1.1	101
7	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.	1.3	99
8	Germline 657del5 mutation in the NBS1 gene in breast cancer patients. International Journal of Cancer, 2003, 106, 379-381.	2.3	80
9	Polymorphism in the P-glycoprotein drug transporter MDR1 gene in colon cancer patients. European Journal of Clinical Pharmacology, 2005, 61, 389-394.	0.8	79
10	CARD15 variants in patients with sporadic Parkinson's disease. Neuroscience Research, 2007, 57, 473-476.	1.0	67
11	Germline CHEK2 mutations and colorectal cancer risk: different effects of a missense and truncating mutations?. European Journal of Human Genetics, 2007, 15, 237-241.	1.4	61
12	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.	1.5	47
13	The 3020insC allele of NOD2 predisposes to early-onset breast cancer. Breast Cancer Research and Treatment, 2005, 89, 91-93.	1.1	47
14	Mutation analysis of MLH1 and MSH2 genes performed by denaturing high-performance liquid chromatography. Journal of Proteomics, 2002, 51, 89-100.	2.4	44
15	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.	1.5	44
16	CDKN2A common variants and their association with melanoma risk: a population-based study. Cancer Research, 2005, 65, 835-9.	0.4	43
17	Inflammatory response gene polymorphisms and their relationship with colorectal cancer risk. BMC Cancer, 2008, 8, 112.	1.1	41
18	Haemochromatosis HFE gene polymorphisms as potential modifiers of hereditary nonpolyposis colorectal cancer risk and onset age. International Journal of Cancer, 2009, 125, 78-83.	2.3	39

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19	MC1R common variants, CDKN2A and their association with melanoma and breast cancer risk. International Journal of Cancer, 2006, 119, 2597-2602.	2.3	38
20	Germline MSH2 and MLH1 mutational spectrum including large rearrangements in HNPCC families from Poland (update study). Clinical Genetics, 2005, 69, 40-47.	1.0	34
21	BRCA1 mutations and colorectal cancer in Poland. Familial Cancer, 2010, 9, 541-544.	0.9	33
22	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.	2.3	33
23	Polymorphisms in nucleotide excision repair genes and susceptibility to colorectal cancer in the Polish population. Molecular Biology Reports, 2015, 42, 755-764.	1.0	32
24	Prevalence of the NOD2 3020insC mutation in aggregations of breast and lung cancer. Breast Cancer Research and Treatment, 2006, 95, 141-145.	1.1	31
25	Germline 657del5 mutation in the NBS1 gene in patients with malignant melanoma of the skin. Melanoma Research, 2003, 13, 365-370.	0.6	30
26	Relationship between acetylation polymorphism and risk of atopic diseases. Clinical Pharmacology and Therapeutics, 1999, 65, 562-569.	2.3	29
27	Nonalcoholic fatty liver disease andHFEgene mutations: A Polish study. World Journal of Gastroenterology, 2010, 16, 2531.	1.4	29
28	Importance of microsatellite instability (MSI) in colorectal cancer: MSI as a diagnostic tool. Annals of Oncology, 2004, 15, iv283-iv284.	0.6	28
29	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.	2.3	28
30	Variant alleles of the CYP1B1 gene are associated with colorectal cancer susceptibility. BMC Cancer, 2010, 10, 420.	1.1	28
31	CHEK2 mutations and HNPCCâ€related colorectal cancer. International Journal of Cancer, 2010, 126, 3005-3009.	2.3	28
32	Fluorescence In Situ Detection of Human Cutaneous Melanoma: Study of Diagnostic Parameters of the Method. Journal of Investigative Dermatology, 2001, 117, 1449-1451.	0.3	26
33	Germline MSH2 and MLH1 mutational spectrum in HNPCC families from Poland and the Baltic States. Journal of Medical Genetics, 2002, 39, 65e-65.	1.5	26
34	The 3020insC Allele of NOD2 Predisposes to Cancers of Multiple Organs. Hereditary Cancer in Clinical Practice, 2005, 3, 59.	0.6	26
35	CDKN2A common variant and multi-organ cancer riskâ€”a population-based study. International Journal of Cancer, 2006, 118, 3180-3182.	2.3	26
36	IGF1 is a modifier of disease risk in hereditary nonâ€polyposis colorectal cancer. International Journal of Cancer, 2008, 123, 1339-1343.	2.3	25

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37	DNA repair gene polymorphisms and risk of early onset colorectal cancer in Lynch syndrome. Cancer Epidemiology, 2012, 36, 183-189.	0.8	25
38	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.	1.4	24
39	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.	2.3	24
40	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.	1.1	23
41	Polymorphism of GSTM1 gene in patients with colorectal cancer and colonic polyps. Experimental and Toxicologic Pathology, 1999, 51, 321-325.	2.1	22
42	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.	0.8	22
43	Analysis of ??+ T cells in peripheral blood of children with perinatal human immunodeficiency virus (HIV) infection. Journal of Clinical Immunology, 1993, 13, 193-203.	2.0	20
44	Optimization of experimental conditions for RNA-based sequencing of MLH1 and MSH2 genes. Human Mutation, 2001, 17, 52-60.	1.1	19
45	Association of MMP8 gene variation with an increased risk of malignant melanoma. Melanoma Research, 2011, 21, 464-468.	0.6	19
46	Long polymerase chain reaction in detection of germline deletions in the von Hippel-Lindau tumour suppressor gene. Human Genetics, 1999, 105, 333-336.	1.8	18
47	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.	0.6	18
48	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.	1.3	17
49	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.	1.4	17
50	NOD2 variants and the risk of malignant melanoma. European Journal of Cancer Prevention, 2005, 14, 143-146.	0.6	16
51	AMPD1 gene mutations are associated with obesity and diabetes in Polish patients with cardiovascular diseases. Journal of Applied Genetics, 2011, 52, 67-76.	1.0	16
52	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.	0.6	14
53	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.	1.3	12
54	Ovarian cancer of endometrioid type as part of the MSH6 gene mutation phenotype. Journal of Human Genetics, 2002, 47, 0529-0531.	1.1	12

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55	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.	0.3	12
56	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.	0.6	11
57	Frequency of mutations related to hereditary haemochromatosis in northwestern Poland. Journal of Applied Genetics, 2008, 49, 105-107.	1.0	10
58	The â^'149C>T SNP within the Î"DNMT3B gene, is not associated with early disease onset in hereditary non-polyposis colorectal cancer. Cancer Letters, 2008, 265, 39-44.	3.2	10
59	CD36 gene polymorphism and plasma sCD36 as the risk factor in higher cholesterolemia. Archives De Pediatrie, 2018, 25, 177-181.	0.4	10
60	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.	1.2	8
61	Nuclear Pedigree Criteria of Suspected HNPCC. Hereditary Cancer in Clinical Practice, 2003, 1, 1.	0.6	8
62	Rarity of germline 1100delC mutation in CHK2 in patients with malignant melanoma of the skin. Melanoma Research, 2004, 14, 121-124.	0.6	8
63	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.	1.2	8
64	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.	0.4	7
65	Frequency and nature of hMSH6 germline mutations in Polish patients with colorectal, endometrial and ovarian cancers. Clinical Genetics, 2006, 70, 68-70.	1.0	6
66	Analysis of HumanCD36Gene 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.	0.3	6
67	DNA and RNA analyses in detection of genetic predisposition to cancer. Hereditary Cancer in Clinical Practice, 2012, 10, 17.	0.6	6
68	Is CD36 gene polymorphism in region encoding lipid-binding domain associated with early onset CAD?. Gene, 2013, 530, 134-137.	1.0	6
69	New <scp>EPCAM</scp> founder deletion in Polish population. Clinical Genetics, 2017, 92, 649-653.	1.0	6
70	A genetic variant in telomerase reverse transcriptase (TERT) modifies cancer risk in Lynch syndrome patients harbouring pathogenic MSH2 variants. Scientific Reports, 2021, 11, 11401.	1.6	6
71	Lynch syndrome mutations shared by the Baltic States and Poland. Clinical Genetics, 2014, 86, 190-193.	1.0	5
72	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.	0.7	5

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73	Cumulative effects of genetic markers and the detection of advanced colorectal neoplasias by population screening. Clinical Genetics, 2015, 88, 234-240.	1.0	5
74	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
75	Nationwide study of clinical and molecular features of hereditary non-polyposis colorectal cancer (HNPCC) in Latvia. Anticancer Research, 2007, 27, 653-8.	0.5	4
76	DNA testing for variants conferring low or moderate increase in the risk of cancer. Hereditary Cancer in Clinical Practice, 2008, 6, 84.	0.6	3
77	Molecular basis of inherited predispositions for tumors.. Acta Biochimica Polonica, 2002, 49, 571-581.	0.3	3
78	Electro-Oculographic and Electroretinographic Studies in HNPCC Gene Mutation Carriers. Ophthalmic Research, 2003, 35, 281-294.	1.0	2
79	Clinical characteristics of tumors derived from colorectal cancer patients who harbor the Tumor Necrosis Factor Î±-1031T/T and NOD2 3020insC polymorphism. Cancer Epidemiology, 2009, 33, 161-163.	0.8	2
80	Hereditary breast cancer. Polish Journal of Pathology, 1998, 49, 59-66.	0.1	2
81	HFE gene mutations in patients with alcoholic liver disease. A prospective study from northwestern Poland. , 2010, 120, 127-31.		2
82	Low-risk Genes and Multi-organ Cancer Risk in the Polish Population. Hereditary Cancer in Clinical Practice, 2006, 4, 52.	0.6	1
83	Some aspects of molecular diagnostics in Lynch syndrome. Hereditary Cancer in Clinical Practice, 2006, 4, 197.	0.6	1
84	DNA and RNA analyses in detection of genetic predisposition to cancer. Hereditary Cancer in Clinical Practice, 2008, 6, 73.	0.6	0
85	MSH2 and MLH1 testing. Hereditary Cancer in Clinical Practice, 2008, 6, 83.	0.6	0
86	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.	0.6	0
87	Molecular Analyses in Diagnosis of High Genetic Predispositions to Malignancies.. Journal of Clinical Biochemistry and Nutrition, 2000, 28, 159-165.	0.6	0