Cezary Cybulski

List of Publications by Year
in descending order

162	9,107	61857	49773
papers	citations	43 h-index	87 g-index

170	170	170	13636
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Validated biomarker assays confirm that <scp>ARID1A</scp> loss is confounded with <scp>MMR</scp> deficiency, <scp>CD8⁺ TIL</scp> infiltration, and provides no independent prognostic value in endometriosisâ€associated ovarian carcinomas. Journal of Pathology, 2022, 256, 388-401.	2.1	15
2	Prostate cancer risk stratification improvement across multiple ancestries with new polygenic hazard score. Prostate Cancer and Prostatic Diseases, 2022, 25, 755-761.	2.0	14
3	Variant Identification in BARD1, PRDM9, RCC1, and RECQL in Patients with Ovarian Cancer by Targeted Next-generation Sequencing of DNA Pools. Cancer Prevention Research, 2022, 15, 151-160.	0.7	2
4	Risk of Second Primary Thyroid Cancer in Women with Breast Cancer. Cancers, 2022, 14, 957.	1.7	5
5	Association of recurrent mutations in BRCA1, BRCA2, RAD51C, PALB2, and CHEK2 with the risk of borderline ovarian tumor. Hereditary Cancer in Clinical Practice, 2022, 20, 11.	0.6	4
6	The impact of oophorectomy on survival from breast cancer in patients with CHEK2 mutations. British Journal of Cancer, 2022, 127, 84-91.	2.9	4
7	Frequency of BRCA1 and BRCA2 mutations in ovarian cancer patients in South-East Poland. Hereditary Cancer in Clinical Practice, 2022, 20, 12.	0.6	0
8	Common Variant in ALDH2 Modifies the Risk of Breast Cancer Among Carriers of the p.K3326* Variant in BRCA2. JCO Precision Oncology, 2022, 6, e2100450.	1.5	1
9	An appraisal of genetic testing for prostate cancer susceptibility. Npj Precision Oncology, 2022, 6, .	2.3	6
10	Population-based targeted sequencing of 54 candidate genes identifiesPALB2as a susceptibility gene for high-grade serous ovarian cancer. Journal of Medical Genetics, 2021, 58, 305-313.	1.5	26
11	Lung Cancer Occurrenceâ€"Correlation with Serum Chromium Levels and Genotypes. Biological Trace Element Research, 2021, 199, 1228-1236.	1.9	13
12	Blood cadmium levels as a marker for early lung cancer detection. Journal of Trace Elements in Medicine and Biology, 2021, 64, 126682.	1.5	28
13	Prevalence of germline TP53 variants among early-onset breast cancer patients from Polish population. Breast Cancer, 2021, 28, 226-235.	1.3	10
14	Trans-ancestry genome-wide association meta-analysis of prostate cancer identifies new susceptibility loci and informs genetic risk prediction. Nature Genetics, 2021, 53, 65-75.	9.4	264
15	Recurrent PALB2 mutations and the risk of cancers of bladder or kidney in Polish population. Hereditary Cancer in Clinical Practice, 2021, 19, 6.	0.6	1
16	Additional SNPs improve risk stratification of a polygenic hazard score for prostate cancer. Prostate Cancer and Prostatic Diseases, 2021, 24, 532-541.	2.0	16
17	Polygenic hazard score is associated with prostate cancer in multi-ethnic populations. Nature Communications, 2021, 12, 1236.	5.8	40
18	Recurrent Mutations in BRCA1, BRCA2, RAD51C, PALB2 and CHEK2 in Polish Patients with Ovarian Cancer. Cancers, 2021, 13, 849.	1.7	13

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19	Survival from breast cancer in women with a BRCA2 mutation by treatment. British Journal of Cancer, 2021, 124, 1524-1532.	2.9	12
20	Serum Selenium Level Predicts 10-Year Survival after Breast Cancer. Nutrients, 2021, 13, 953.	1.7	14
21	KLK3 SNPâ€“SNP interactions for prediction of prostate cancer aggressiveness. Scientific Reports, 2021, 11, 9264.	1.6	5
22	PALB2 mutations and prostate cancer risk and survival. British Journal of Cancer, 2021, 125, 569-575.	2.9	18
23	Blood Arsenic Levels as a Marker of Breast Cancer Risk among BRCA1 Carriers. Cancers, 2021, 13, 3345.	1.7	6
24	Serum Selenium Level and 10-Year Survival after Melanoma. Biomedicines, 2021, 9, 991.	1.4	8
25	Low Blood-As Levels and Selected Genotypes Appears to Be Promising Biomarkers for Occurrence of Colorectal Cancer in Women. Biomedicines, 2021, 9, 1105.	1.4	0
26	Genetic predisposition to male breast cancer in Poland. BMC Cancer, 2021, 21, 975.	1.1	7
27	Influence of the Levels of Arsenic, Cadmium, Mercury and Lead on Overall Survival in Lung Cancer. Biomolecules, 2021, 11, 1160.	1.8	23
28	Survival of bladder or renal cancer in patients with CHEK2 mutations. PLoS ONE, 2021, 16, e0257132.	1.1	1
29	BRCA1 and BRCA2 mutations in ovarian cancer patients from Belarus: update. Hereditary Cancer in Clinical Practice, 2021, 19, 13.	0.6	8
30	Do BARD1 Mutations Confer an Elevated Risk of Prostate Cancer?. Cancers, 2021, 13, 5464.	1.7	1
31	Blood Copper Levels and the Occurrence of Colorectal Cancer in Poland. Biomedicines, 2021, 9, 1628.	1.4	19
32	Blood arsenic levels and the risk of familial breast cancer in Poland. International Journal of Cancer, 2020, 146, 2721-2727.	2.3	18
33	BRCA1 promoter methylation in peripheral blood is associated with the risk of tripleâ€negative breast cancer. International Journal of Cancer, 2020, 146, 1293-1298.	2.3	33
34	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. Nature Genetics, 2020, 52, 56-73.	9.4	120
35	Constitutional variants in POT1, TERF2IP, and ACD genes in patients with melanoma in the Polish population. European Journal of Cancer Prevention, 2020, 29, 511-519.	0.6	3
36	Polymorphisms in MMP-1, MMP-2, MMP-7, MMP-13 and MT2A do not contribute to breast, lung and colon cancer risk in polish population. Hereditary Cancer in Clinical Practice, 2020, 18, 16.	0.6	14

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37	Prevalence of Recurrent Mutations Predisposing to Breast Cancer in Early-Onset Breast Cancer Patients from Poland. Cancers, 2020, 12, 2321.	1.7	11
38	Mutations in ATM , NBN and BRCA2 predispose to aggressive prostate cancer in Poland. International Journal of Cancer, 2020, 147, 2793-2800.	2.3	27
39	CA125 and Ovarian Cancer: A Comprehensive Review. Cancers, 2020, 12, 3730.	1.7	174
40	The CHEK2 Variant C.349A>G Is Associated with Prostate Cancer Risk and Carriers Share a Common Ancestor. Cancers, 2020, 12, 3254.	1.7	16
41	The effect of sample size on polygenic hazard models for prostate cancer. European Journal of Human Genetics, 2020, 28, 1467-1475.	1.4	14
42	Development and Validation of the Gene Expression Predictor of High-grade Serous Ovarian Carcinoma Molecular SubTYPE (PrOTYPE). Clinical Cancer Research, 2020, 26, 5411-5423.	3.2	43
43	A Genetic Risk Score to Personalize Prostate Cancer Screening, Applied to Population Data. Cancer Epidemiology Biomarkers and Prevention, 2020, 29, 1731-1738.	1.1	27
44	Transcriptomeâ€wide association study of breast cancer risk by estrogenâ€receptor status. Genetic Epidemiology, 2020, 44, 442-468.	0.6	32
45	Influence of the selenium level on overall survival in lung cancer. Journal of Trace Elements in Medicine and Biology, 2019, 56, 46-51.	1.5	25
46	Allelic modification of breast cancer risk in women with an NBN mutation. Breast Cancer Research and Treatment, 2019, 178, 427-431.	1.1	6
47	Sex specific associations in genome wide association analysis of renal cell carcinoma. European Journal of Human Genetics, 2019, 27, 1589-1598.	1.4	27
48	Inherited Variants in BLM and the Risk and Clinical Characteristics of Breast Cancer. Cancers, 2019, 11, 1548.	1.7	11
49	Inherited variants in XRCC2 and the risk of breast cancer. Breast Cancer Research and Treatment, 2019, 178, 657-663.	1.1	13
50	Interim Results from the IMPACT Study: Evidence for Prostate-specific Antigen Screening in BRCA2 Mutation Carriers. European Urology, 2019, 76, 831-842.	0.9	148
51	A comparison of ovarian cancer mortality in women with BRCA1 mutations undergoing annual ultrasound screening or preventive oophorectomy. Gynecologic Oncology, 2019, 155, 270-274.	0.6	15
52	Shared heritability and functional enrichment across six solid cancers. Nature Communications, 2019, 10, 431.	5.8	88
53	The Relationship between the HLA-G Polymorphism and sHLA-G Levels in Parental Pairs with High-Risk Pregnancy. International Journal of Environmental Research and Public Health, 2019, 16, 1546.	1.2	7
54	The spectrum of mutations predisposing to familial breast cancer in Poland. International Journal of Cancer, 2019, 145, 3311-3320.	2.3	39

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55	BARD1 is a Low/Moderate Breast Cancer Risk Gene: Evidence Based on an Association Study of the Central European p.Q564X Recurrent Mutation. Cancers, 2019, 11, 740.	1.7	25
56	Molecular Analysis of HLA-G in Women with High-Risk Pregnancy and Their Partners with Regard to Possible Complications. International Journal of Environmental Research and Public Health, 2019, 16, 982.	1.2	23
57	CHEK2 Alleles Predispose to Renal Cancer in Poland. JAMA Oncology, 2019, 5, 576.	3.4	8
58	Serum selenium level and cancer risk: a nested case-control study. Hereditary Cancer in Clinical Practice, 2019, 17, 33.	0.6	15
59	The influence of obesity-related factors in the etiology of renal cell carcinomaâ€"A mendelian randomization study. PLoS Medicine, 2019, 16, e1002724.	3.9	59
60	Age-specific risks of incident, contralateral and ipsilateral breast cancer among 1776 Polish BRCA1 mutation carriers. Breast Cancer Research and Treatment, 2019, 174, 769-774.	1.1	7
61	Circulating Metabolic Biomarkers of Screen-Detected Prostate Cancer in the ProtecT Study. Cancer Epidemiology Biomarkers and Prevention, 2019, 28, 208-216.	1.1	21
62	Founder Mutations for Early Onset Melanoma as Revealed by Whole Exome Sequencing Suggests That This is Not Associated with the Increasing Incidence of Melanoma in Poland. Cancer Research and Treatment, 2019, 51, 337-344.	1.3	8
63	Inherited NBN Mutations and Prostate Cancer Risk and Survival. Cancer Research and Treatment, 2019, 51, 1180-1187.	1.3	21
64	Predictors of survival for breast cancer patients with a BRCA1 mutation. Breast Cancer Research and Treatment, 2018, 168, 513-521.	1.1	20
65	Prostate-specific antigen velocity in a prospective prostate cancer screening study of men with genetic predisposition. British Journal of Cancer, 2018, 118, 266-276.	2.9	12
66	Polygenic hazard score to guide screening for aggressive prostate cancer: development and validation in large scale cohorts. BMJ: British Medical Journal, 2018, 360, j5757.	2.4	153
67	Germline variation at 8q24 and prostate cancer risk in men of European ancestry. Nature Communications, 2018, 9, 4616.	5.8	43
68	BRCA1/2 mutations are not a common cause of malignant melanoma in the Polish population. PLoS ONE, 2018, 13, e0204768.	1.1	6
69	Reply to â€˜Mutations in RECQL are not associated with breast cancer risk in an Australian populationâ€™. Nature Genetics, 2018, 50, 1348-1349.	9.4	2
70	Testing Ashkenazi Jewish Women for Mutations Predisposing to Breast Cancer in Genes Other Than BRCA1 and BRCA2. JAMA Oncology, 2018, 4, 1012.	3.4	3
71	Association of p16 expression with prognosis varies across ovarian carcinoma histotypes: an Ovarian Tumor Tissue Analysis consortium study. Journal of Pathology: Clinical Research, 2018, 4, 250-261.	1.3	70
72	Association of zinc level and polymorphism in MMP-7 gene with prostate cancer in Polish population. PLoS ONE, 2018, 13, e0201065.	1.1	30

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73	Variants in genes encoding small GTPases and association with epithelial ovarian cancer susceptibility. PLoS ONE, 2018, 13, e0197561.	1.1	9
74	rs495139 in the TYMS-ENOSF1 Region and Risk of Ovarian Carcinoma of Mucinous Histology. International Journal of Molecular Sciences, 2018, 19, 2473.	1.8	3
75	AA9int: SNP interaction pattern search using non-hierarchical additive model set. Bioinformatics, 2018, 34, 4141-4150.	1.8	3
76	Association analyses of more than 140,000 men identify 63 new prostate cancer susceptibility loci. Nature Genetics, 2018, 50, 928-936.	9.4	652
77	Fine-mapping of prostate cancer susceptibility loci in a large meta-analysis identifies candidate causal variants. Nature Communications, 2018, 9, 2256.	5.8	88
78	SNP interaction pattern identifier (SIPI): an intensive search for SNPâ€“SNP interaction patterns. Bioinformatics, 2017, 33, 822-833.	1.8	11
79	Serum 25(OH)D concentration, common variants of the VDR gene and lung cancer occurrence. International Journal of Cancer, 2017, 141, 336-341.	2.3	16
80	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724.	5.8	106
81	BRCA1/2â€negative hereditary tripleâ€negative breast cancers exhibit BRCAness. International Journal of Cancer, 2017, 140, 1545-1550.	2.3	9
82	Screening with magnetic resonance imaging, mammography and ultrasound in women at average and intermediate risk of breast cancer. Hereditary Cancer in Clinical Practice, 2017, 15, 4.	0.6	17
83	Investigating the possible causal role of coffee consumption with prostate cancer risk and progression using Mendelian randomization analysis. International Journal of Cancer, 2017, 140, 322-328.	2.3	17
84	Alcohol consumption and prostate cancer incidence and progression: A Mendelian randomisation study. International Journal of Cancer, 2017, 140, 75-85.	2.3	28
85	Body mass index and breast cancer survival: a Mendelian randomization analysis. International Journal of Epidemiology, 2017, 46, 1814-1822.	0.9	45
86	The 30 kb deletion in the APOBEC3 cluster decreases APOBEC3A and APOBEC3B expression and creates a transcriptionally active hybrid gene but does not associate with breast cancer in the European population. Oncotarget, 2017, 8, 76357-76374.	0.8	26
87	The Prevalence of Founder Mutations among Individuals from Families with Familial Pancreatic Cancer Syndrome. Cancer Research and Treatment, 2017, 49, 430-436.	1.3	19
88	Exome genotyping arrays to identify rare and low frequency variants associated with epithelial ovarian cancer risk. Human Molecular Genetics, 2016, 25, 3600-3612.	1.4	17
89	Do founder mutations characteristic of some cancer sites also predispose to pancreatic cancer?. International Journal of Cancer, 2016, 139, 601-606.	2.3	16
90	PALB2,CHEK2andATMrare variants and cancer risk: data from COGS. Journal of Medical Genetics, 2016, 53, 800-811.	1.5	174

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91	Polyunsaturated fatty acids and prostate cancer risk: a Mendelian randomisation analysis from the PRACTICAL consortium. British Journal of Cancer, 2016, 115, 624-631.	2.9	23
92	Assessing the role of insulinâ€like growth factors and binding proteins in prostate cancer using Mendelian randomization: Genetic variants as instruments for circulating levels. International Journal of Cancer, 2016, 139, 1520-1533.	2.3	26
93	Blood lipids and prostate cancer: a Mendelian randomization analysis. Cancer Medicine, 2016, 5, 1125-1136.	1.3	68
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.	7.7	157
95	Inheritance of deleterious mutations at both BRCA1 and BRCA2 in an international sample of 32,295 women. Breast Cancer Research, 2016, 18, 112.	2.2	42
96	Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breastâ€“ovarian cancer susceptibility locus. Nature Communications, 2016, 7, 12675.	5.8	78
97	Atlas of prostate cancer heritability in European and African-American men pinpoints tissue-specific regulation. Nature Communications, 2016, 7, 10979.	5.8	50
98	Pubertal development and prostate cancer risk: Mendelian randomization study in a population-based cohort. BMC Medicine, 2016, 14, 66.	2.3	42
99	Assessment of variation in immunosuppressive pathway genes reveals TGFBR2 to be associated with risk of clear cell ovarian cancer. Oncotarget, 2016, 7, 69097-69110.	0.8	5
100	Pathological complete response after cisplatin neoadjuvant therapy is associated with the downregulation of DNA repair genes in BRCA1-associated triple-negative breast cancers. Oncotarget, 2016, 7, 68662-68673.	0.8	13
101	Epithelialâ€Mesenchymal Transition (EMT) Gene Variants and Epithelial Ovarian Cancer (EOC) Risk. Genetic Epidemiology, 2015, 39, 689-697.	0.6	22
102	Prediction of individual genetic risk to prostate cancer using a polygenic score. Prostate, 2015, 75, 1467-1474.	1.2	54
103	Review Selenium as aÂmarker of cancer risk and of selection for control examinations in surveillance. Wspolczesna Onkologia, 2015, 1A, 60-61.	0.7	4
104	Common Genetic Variation In Cellular Transport Genes and Epithelial Ovarian Cancer (EOC) Risk. PLoS ONE, 2015, 10, e0128106.	1.1	44
105	Prevalence of Germline Mutations in Genes Engaged in DNA Damage Repair by Homologous Recombination in Patients with Triple-Negative and Hereditary Non-Triple-Negative Breast Cancers. PLoS ONE, 2015, 10, e0130393.	1.1	22
106	Prospective evaluation of alcohol consumption and the risk of breast cancer in BRCA1 and BRCA2 mutation carriers. Breast Cancer Research and Treatment, 2015, 151, 435-441.	1.1	12
107	A Large-Scale Analysis of Genetic Variants within Putative miRNA Binding Sites in Prostate Cancer. Cancer Discovery, 2015, 5, 368-379.	7.7	56
108	Identification of six new susceptibility loci for invasive epithelial ovarian cancer. Nature Genetics, 2015, 47, 164-171.	9.4	221

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109	CHEK2mutations and the risk of papillary thyroid cancer. International Journal of Cancer, 2015, 137, 548-552.	2.3	97
110	Network-Based Integration of GWAS and Gene Expression Identifies a HOX-Centric Network Associated with Serous Ovarian Cancer Risk. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1574-1584.	1.1	28
111	Genome-wide Analysis Identifies Novel Loci Associated with Ovarian Cancer Outcomes: Findings from the Ovarian Cancer Association Consortium. Clinical Cancer Research, 2015, 21, 5264-5276.	3.2	33
112	Germline RECQL mutations are associated with breast cancer susceptibility. Nature Genetics, 2015, 47, 643-646.	9.4	168
113	Risk Analysis of Prostate Cancer in PRACTICAL, a Multinational Consortium, Using 25 Known Prostate Cancer Susceptibility Loci. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1121-1129.	1.1	56
114	Evaluating the ovarian cancer gonadotropin hypothesis: A candidate gene study. Gynecologic Oncology, 2015, 136, 542-548.	0.6	15
115	Association of Type and Location ofBRCA1andBRCA2Mutations With Risk of Breast and Ovarian Cancer. JAMA - Journal of the American Medical Association, 2015, 313, 1347.	3.8	390
116	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
117	Clinical outcomes in women with breast cancer and a PALB2 mutation: a prospective cohort analysis. Lancet Oncology, The, 2015, 16, 638-644.	5.1	137
118	Cis-eQTL analysis and functional validation of candidate susceptibility genes for high-grade serous ovarian cancer. Nature Communications, 2015, 6, 8234.	5.8	63
119	Common variants at theCHEK2gene locus and risk of epithelial ovarian cancer. Carcinogenesis, 2015, 36, 1341-1353.	1.3	24
120	The effects of height and BMI on prostate cancer incidence and mortality: a Mendelian randomization study in 20,848 cases and 20,214 controls from the PRACTICAL consortium. Cancer Causes and Control, 2015, 26, 1603-1616.	0.8	77
121	Multiple novel prostate cancer susceptibility signals identified by fine-mapping of known risk loci among Europeans. Human Molecular Genetics, 2015, 24, 5589-5602.	1.4	67
122	Genome-Wide Association Study of Prostate Cancerâ€“Specific Survival. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 1796-1800.	1.1	27
123	Factors influencing ovulation and the risk of ovarian cancer in <scp>BRCA1</scp> and <scp>BRCA2</scp> mutation carriers. International Journal of Cancer, 2015, 137, 1136-1146.	2.3	56
124	Candidate Genetic Modifiers for Breast and Ovarian Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 308-316.	1.1	22
125	RECQL: a DNA helicase in breast cancer. Oncotarget, 2015, 6, 26558-26559.	0.8	10
126	Common Genetic Variation in Circadian Rhythm Genes and Risk of Epithelial Ovarian Cancer (EOC). Journal of Genetics and Genome Research, 2015, 2, .	0.3	25

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127	DNA Glycosylases Involved in Base Excision Repair May Be Associated with Cancer Risk in BRCA1 and BRCA2 Mutation Carriers. PLoS Genetics, 2014, 10, e1004256.	1.5	47
128	Fine-Mapping the HOXB Region Detects Common Variants Tagging a Rare Coding Allele: Evidence for Synthetic Association in Prostate Cancer. PLoS Genetics, 2014, 10, e1004129.	1.5	34
129	Variation in NF-Î¹B Signaling Pathways and Survival in Invasive Epithelial Ovarian Cancer. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 1421-1427.	1.1	13
130	Multiple primary cancers as a guide to heritability. International Journal of Cancer, 2014, 135, 1756-1763.	2.3	55
131	Targeted Prostate Cancer Screening in BRCA1 and BRCA2 Mutation Carriers: Results from the Initial Screening Round of the IMPACT Study. European Urology, 2014, 66, 489-499.	0.9	195
132	Impact of Oophorectomy on Cancer Incidence and Mortality in Women With a BRCA1 or BRCA2 Mutation. Journal of Clinical Oncology, 2014, 32, 1547-1553.	0.8	523
133	First recurrent large genomic rearrangement in the BRCA1 gene found in Poland. Cancer Epidemiology, 2014, 38, 382-385.	0.8	1
134	A meta-analysis of 87,040 individuals identifies 23 new susceptibility loci for prostate cancer. Nature Genetics, 2014, 46, 1103-1109.	9.4	408
135	Prevalence of the E318K and V320I MITF germline mutations in Polish cancer patients and multiorgan cancer risk-a population-based study. Cancer Genetics, 2014, 207, 128-132.	0.2	23
136	Common variants of xeroderma pigmentosum genes and prostate cancer risk. Gene, 2014, 546, 156-161.	1.0	23
137	Population distribution and ancestry of the cancer protective MDM2 SNP285 (rs117039649). Oncotarget, 2014, 5, 8223-8234.	0.8	22
138	GWAS meta-analysis and replication identifies three new susceptibility loci for ovarian cancer. Nature Genetics, 2013, 45, 362-370.	9.4	326
139	The G84E mutation in the HOXB13 gene is associated with an increased risk of prostate cancer in Poland. Prostate, 2013, 73, 542-548.	1.2	31
140	A common nonsense mutation of the BLM gene and prostate cancer risk and survival. Gene, 2013, 532, 173-176.	1.0	24
141	Multiple independent variants at the TERT locus are associated with telomere length and risks of breast and ovarian cancer. Nature Genetics, 2013, 45, 371-384.	9.4	493
142	From Phenotype to Genotype: A New Twist on Identifying Genes Responsible for Inherited Hearing Loss. Human Mutation, 2013, 34, v-v.	1.1	2
143	The risk of gastric cancer in carriers of CHEK2 mutations. Familial Cancer, 2013, 12, 473-478.	0.9	46
144	A meta-analysis of genome-wide association studies to identify prostate cancer susceptibility loci associated with aggressive and non-aggressive disease. Human Molecular Genetics, 2013, 22, 408-415.	1.4	118

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145	Genotyping by Induced FÃ¶rster Resonance Energy Transfer (iFRET) Mechanism and Simultaneous Mutation Scanning. Human Mutation, 2013, 34, n/a-n/a.	1.1	2
146	Epigenetic analysis leads to identification of HNF1B as a subtype-specific susceptibility gene for ovarian cancer. Nature Communications, 2013, 4, 1628.	5.8	144
147	A Low Selenium Level Is Associated with Lung and Laryngeal Cancers. PLoS ONE, 2013, 8, e59051.	1.1	46
148	Risk of Breast Cancer in Women With a CHEK2 Mutation With and Without a Family History of Breast Cancer. Journal of Clinical Oncology, 2011, 29, 3747-3752.	0.8	207
149	Genetic Variation at 9p22.2 and Ovarian Cancer Risk for BRCA1 and BRCA2 Mutation Carriers. Journal of the National Cancer Institute, 2011, 103, 105-116.	3.0	40
150	Constitutional CHEK2 mutations are associated with a decreased risk of lung and laryngeal cancers. Carcinogenesis, 2008, 29, 762-765.	1.3	41
151	BRCA1 mutations and prostate cancer in Poland. European Journal of Cancer Prevention, 2008, 17, 62-66.	0.6	33
152	Epistatic Relationship between the Cancer Susceptibility Genes CHEK2 and p27. Cancer Epidemiology Biomarkers and Prevention, 2007, 16, 572-576.	1.1	21
153	DNA Variation in MSR1, RNASEL and E-Cadherin Genes and Prostate Cancer in Poland. Urologia Internationalis, 2007, 79, 44-49.	0.6	20
154	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
155	A deletion in CHEK2 of 5,395Âbp predisposes to breast cancer in Poland. Breast Cancer Research and Treatment, 2007, 102, 119-122.	1.1	102
156	Population Screening of CHEK2 Mutations in Poland. Hereditary Cancer in Clinical Practice, 2006, 4, 57.	0.6	2
157	CHEK2-Positive Breast Cancers in Young Polish Women. Clinical Cancer Research, 2006, 12, 4832-4835.	3.2	32
158	A Novel Founder CHEK2 Mutation is Associated with Increased Prostate Cancer Risk: Table 1. Cancer Research, 2004, 64, 2677-2679.	0.4	137
159	A high proportion of founderBRCA1mutations in Polish breast cancer families. International Journal of Cancer, 2004, 110, 683-686.	2.3	170
160	Mutations in the von Hippel-Lindau Tumour Suppressor Gene in Central Nervous System Hemangioblastomas. Hereditary Cancer in Clinical Practice, 2004, 2, 93.	0.6	2
161	Germline 657del5 mutation in the NBS1 gene in breast cancer patients. International Journal of Cancer, 2003, 106, 379-381.	2.3	80
162	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