

Fine-mapping of the HNF1B multicancer locus identifies endometrial cancer risk

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1. <i>Scientific Reports</i> , 2015, 5, 17369.	1.6	35
2	Candidate locus analysis of the TERT/CLPTM1L cancer risk region on chromosome 5p15 identifies multiple independent variants associated with endometrial cancer risk. <i>Human Genetics</i> , 2015, 134, 231-245.	1.8	34
3	Limited evidence that cancer susceptibility regions are preferential targets for somatic mutation. <i>Genome Biology</i> , 2015, 16, 193.	3.8	19
4	Comprehensive genetic assessment of the ESR1 locus identifies a risk region for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2015, 22, 851-861.	1.6	25
5	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016, 48, 667-674.	9.4	77
6	Genetic Risk Score Mendelian Randomization Shows that Obesity Measured as Body Mass Index, but not Waist:Hip Ratio, Is Causal for Endometrial Cancer. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2016, 25, 1503-1510.	1.1	64
8	A Common Variant at the 14q32 Endometrial Cancer Risk Locus Activates AKT1 through YY1 Binding. <i>American Journal of Human Genetics</i> , 2016, 98, 1159-1169.	2.6	32
9	CYP19A1 fine-mapping and Mendelian randomization: estradiol is causal for endometrial cancer. <i>Endocrine-Related Cancer</i> , 2016, 23, 77-91.	1.6	62
10	Expression, Epigenetic and Genetic Changes of HNF1B in Endometrial Lesions. <i>Pathology and Oncology Research</i> , 2016, 22, 523-530.	0.9	19
11	GWAS meta-analysis of 16 852 women identifies new susceptibility locus for endometrial cancer. <i>Human Molecular Genetics</i> , 2016, 25, ddw092.	1.4	19
12	Diabetes mellitus and gynecologic cancer: molecular mechanisms, epidemiological, clinical and prognostic perspectives. <i>Archives of Gynecology and Obstetrics</i> , 2016, 293, 239-246.	0.8	50
13	Association Between Telomere Length and Risk of Cancer and Non-Neoplastic Diseases. <i>JAMA Oncology</i> , 2017, 3, 636.	3.4	376
14	Endometrial cancer gene panels: clinical diagnostic vs research germline DNA testing. <i>Modern Pathology</i> , 2017, 30, 1048-1068.	2.9	37
15	Current strategies in the diagnosis of endometrial cancer. <i>Archives of Gynecology and Obstetrics</i> , 2017, 296, 5-14.	0.8	12
17	Family history of cancer predicts endometrial cancer risk independently of Lynch Syndrome: Implications for genetic counselling. <i>Gynecologic Oncology</i> , 2017, 147, 381-387.	0.6	30
18	Pathology and Molecular Pathology of Uterine and Ovarian Cancers. , 2017, , 247-278.		0
19	The OncoArray Consortium: A Network for Understanding the Genetic Architecture of Common Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 126-135.	1.1	278
20	Genetic overlap between endometriosis and endometrial cancer: evidence from cross-disease genetic correlation and GWAS meta-analyses. <i>Cancer Medicine</i> , 2018, 7, 1978-1987.	1.3	62

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21	Genome-wide analysis of PDX1 target genes in human pancreatic progenitors. <i>Molecular Metabolism</i> , 2018, 9, 57-68.	3.0	67
22	Genome-wide meta-analysis identifies five new susceptibility loci for pancreatic cancer. <i>Nature Communications</i> , 2018, 9, 556.	5.8	188
23	Common Genetic Variation and Susceptibility to Ovarian Cancer: Current Insights and Future Directions. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018, 27, 395-404.	1.1	33
24	Risk and prognostic factors for endometrial carcinoma after diagnosis of breast or Lynch-associated cancersâ€”A population-based analysis. <i>Cancer Medicine</i> , 2018, 7, 6411-6422.	1.3	9
25	Cumulative evidence for relationships between multiple variants of HNF1B and the risk of prostate and endometrial cancers. <i>BMC Medical Genetics</i> , 2018, 19, 128.	2.1	5
26	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018, 9, 3166.	5.8	178
27	Genome-Wide Association Studies of Endometrial Cancer: Latest Developments and Future Directions. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019, 28, 1095-1102.	1.1	32
28	Influence of IL1R2 polymorphisms on endometrial cancer susceptibility in the Chinese Han population. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e650.	0.6	3
29	Risk of postmenopausal hormone therapy and patient history factors for the survival rate in women with endometrial carcinoma. <i>Archives of Gynecology and Obstetrics</i> , 2020, 301, 289-294.	0.8	5
30	Germline genetic variation in prostate susceptibility does not predict outcomes in the chemoprevention trials PCPT and SELECT. <i>Prostate Cancer and Prostatic Diseases</i> , 2020, 23, 333-342.	2.0	10
31	Analysis of expression, epigenetic, and genetic changes of HNF1B in 130 kidney tumours. <i>Scientific Reports</i> , 2020, 10, 17151.	1.6	5
32	HNF1B, EZH2 and ECI2 in prostate carcinoma. Molecular, immunohistochemical and clinico-pathological study. <i>Scientific Reports</i> , 2020, 10, 14365.	1.6	6
33	Cross-Cancer Genome-Wide Association Study of Endometrial Cancer and Epithelial Ovarian Cancer Identifies Genetic Risk Regions Associated with Risk of Both Cancers. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2021, 30, 217-228.	1.1	12
34	Expression, Epigenetic, and Genetic Changes of HNF1B in Colorectal Lesions: an Analysis of 145 Cases. <i>Pathology and Oncology Research</i> , 2020, 26, 2337-2350.	0.9	7
35	Hepatocyte nuclear factor 1 beta: A perspective in cancer. <i>Cancer Medicine</i> , 2021, 10, 1791-1804.	1.3	15
36	A Review of Functional Characterization of Single Amino Acid Change Mutations in HNF Transcription Factors in MODY Pathogenesis. <i>Protein Journal</i> , 2021, 40, 348-360.	0.7	11
37	A common variant within the HNF1B gene is associated with overall survival of multiple myeloma patients: Results from the IMMEnSE consortium and meta-analysis. <i>Oncotarget</i> , 2016, 7, 59029-59048.	0.8	16
38	Network-directed cis-mediator analysis of normal prostate tissue expression profiles reveals downstream regulatory associations of prostate cancer susceptibility loci. <i>Oncotarget</i> , 2017, 8, 85896-85908.	0.8	2

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39	Endometrial Cancer Among Asian Americans. , 2016, , 219-231.		0
40	Uterine Clear Cell Carcinoma. Molecular Pathology Library, 2017, , 123-142.	0.1	0
41	The prospect of discovering new biomarkers for ovarian cancer based on current knowledge of χ^2 susceptibility loci and genetic variation (Review). International Journal of Molecular Medicine, 2019, 44, 1599-1608.	1.8	2
43	Lung cancer A549 cells suppressed with overexpressed HNF1B or PCDHA13 inhibited PI3K/AKT phosphorylation. Translational Cancer Research, 2020, 9, 3819-3827.	0.4	1
44	Hereditary Endometrial Cancers. , 2020, , 77-95.		1
45	Haplotypes of single cancer driver genes and their local ancestry in a highly admixed long-lived population of Northeast Brazil. Genetics and Molecular Biology, 2022, 45, e20210172.	0.6	1
46	10 Years of GWAS discovery in endometrial cancer: Aetiology, function and translation. EBioMedicine, 2022, 77, 103895.	2.7	11
47	The landscape of GWAS validation; systematic review identifying 309 validated non-coding variants across 130 human diseases. BMC Medical Genomics, 2022, 15, 74.	0.7	28