

CITATION REPORT

List of articles citing

Meta-analysis of genome-wide association studies identifies common susceptibility polymorphisms for colorectal and endometrial cancer near SH2B3 and TSHZ1

DOI: 10.1038/srep17369
Scientific Reports, 2015, 5, 17369.

Source: <https://exaly.com/paper-pdf/60389980/citation-report.pdf>

Version: 2024-04-28

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#	Paper	IF	Citations
33	Five endometrial cancer risk loci identified through genome-wide association analysis. <i>Nature Genetics</i> , 2016 , 48, 667-674	36.3	56
32	Endometrial cancer gene panels: clinical diagnostic vs research germline DNA testing. <i>Modern Pathology</i> , 2017 , 30, 1048-1068	9.8	20
31	Association between MDR1 C3435T polymorphism and colorectal cancer risk: A meta-analysis. <i>Medicine (United States)</i> , 2017 , 96, e9428	1.8	5
30	Genome-wide association study of offspring birth weight in 86 577 women identifies five novel loci and highlights maternal genetic effects that are independent of fetal genetics. <i>Human Molecular Genetics</i> , 2018 , 27, 742-756	5.6	98
29	Identification of Pleiotropic Cancer Susceptibility Variants from Genome-Wide Association Studies Reveals Functional Characteristics. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2018 , 27, 75-85	4	16
28	Genetic susceptibility markers for a breast-colorectal cancer phenotype: Exploratory results from genome-wide association studies. <i>PLoS ONE</i> , 2018 , 13, e0196245	3.7	2
27	Telomerase Reverse Transcriptase Polymorphism rs2736100: A Balancing Act between Cancer and Non-Cancer Disease, a Meta-Analysis. <i>Frontiers in Medicine</i> , 2018 , 5, 41	4.9	15
26	Identification of nine new susceptibility loci for endometrial cancer. <i>Nature Communications</i> , 2018 , 9, 3166	17.4	70
25	Linear isoforms of the long noncoding RNA CDKN2B-AS1 regulate the c-myc-enhancer binding factor RBMS1. <i>European Journal of Human Genetics</i> , 2019 , 27, 80-89	5.3	26
24	Exome-Wide Rare Variant Analysis From the DiscovEHR Study Identifies Novel Candidate Predisposition Genes for Endometrial Cancer. <i>Frontiers in Oncology</i> , 2019 , 9, 574	5.3	8
23	Genome-Wide Association Studies of Endometrial Cancer: Latest Developments and Future Directions. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2019 , 28, 1095-1102	4	21
22	The mRNA-bound Proteome of : Novel Genetic Insight into an Ancient Parasite. <i>Molecular and Cellular Proteomics</i> , 2019 , 18, 1271-1284	7.6	21
21	Linkage analysis revealed risk loci on 6p21 and 18p11.2-q11.2 in familial colon and rectal cancer, respectively. <i>European Journal of Human Genetics</i> , 2019 , 27, 1286-1295	5.3	1
20	Association between dietary fiber and endometrial cancer: a meta-analysis. <i>Nutrition and Cancer</i> , 2020 , 72, 959-967	2.8	2
19	Current and future approaches to screening for endometrial cancer. <i>Best Practice and Research in Clinical Obstetrics and Gynaecology</i> , 2020 , 65, 79-97	4.6	14
18	An in silico approach to identify and prioritize miRNAs target sites polymorphisms in colorectal cancer and obesity. <i>Cancer Medicine</i> , 2020 , 9, 9511-9528	4.8	3
17	Genetic Susceptibility to Endometrial Cancer: Risk Factors and Clinical Management. <i>Cancers</i> , 2020 , 12,	6.6	12

16	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	4	7
15	Association between genetic polymorphisms and endometrial cancer risk: a systematic review. <i>Journal of Medical Genetics</i> , 2020 , 57, 591-600	5.8	15
14	Cross-cancer evaluation of polygenic risk scores for 16 cancer types in two large cohorts. <i>Nature Communications</i> , 2021 , 12, 970	17.4	13
13	Differential genetic influences over colorectal cancer risk and gene expression in large bowel mucosa. <i>International Journal of Cancer</i> , 2021 , 149, 1100-1108	7.5	2
12	Genetic polymorphisms in gynecologic cancers. 2021 , 169-193		
11	The Colon Cancer Family Registry Cohort. 2018 , 427-459		1
10	Cross-Cancer Evaluation of Polygenic Risk Scores for 17 Cancer Types in Two Large Cohorts.		2
9	Systematic classification of shared components of genetic risk for common human diseases.		1
8	Genome-wide association study of offspring birth weight in 86,577 women highlights maternal genetic effects that are independent of fetal genetics.		
7	The mRNA-bound proteome of <i>Leishmania mexicana</i> : novel genetic insight into an ancient parasite.		
6	Cross-cancer genome-wide association study of endometrial cancer and epithelial ovarian cancer identifies genetic risk regions associated with risk of both cancers.		
5	Hereditary Endometrial Cancers. 2020 , 77-95		
4	10 Years of GWAS discovery in endometrial cancer: Aetiology, function and translation.. <i>EBioMedicine</i> , 2022 , 77, 103895	8.8	2
3	Data_Sheet_1.xlsx. 2019 ,		
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1	Control of <i>Mycobacterium avium</i> subsp. <i>paratuberculosis</i> load within infected bovine monocyte-derived macrophages is associated with host genetics. 14,		0