

Emily Webb

List of Publications by Year in descending order

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14
papers

3,753
citations

686830

13
h-index

1058022

14
g-index

14
all docs

14
docs citations

14
times ranked

4433
citing authors

#	ARTICLE	IF	CITATIONS
1	Comprehensive genetic analysis of seven large families with mismatch repair proficient colorectal cancer. <i>Genes Chromosomes and Cancer</i> , 2010, 49, 539-548.	1.5	17
2	The colorectal cancer risk at 18q21 is caused by a novel variant altering <i>SMAD7</i> expression. <i>Genome Research</i> , 2009, 19, 987-993.	2.4	85
3	A genome-wide scan of 10 ⁶ gene-centric variants and colorectal cancer risk. <i>European Journal of Human Genetics</i> , 2009, 17, 1507-1514.	1.4	12
4	A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. <i>Nature Genetics</i> , 2008, 40, 623-630.	9.4	514
5	Genome-wide association scan identifies a colorectal cancer susceptibility locus on 11q23 and replicates risk loci at 8q24 and 18q21. <i>Nature Genetics</i> , 2008, 40, 631-637.	9.4	542
6	Common genetic variants at the CRAC1 (HMPS) locus on chromosome 15q13.3 influence colorectal cancer risk. <i>Nature Genetics</i> , 2008, 40, 26-28.	9.4	277
7	A genome-wide association study identifies six susceptibility loci for chronic lymphocytic leukemia. <i>Nature Genetics</i> , 2008, 40, 1204-1210.	9.4	329
8	Meta-analysis of genome-wide association data identifies four new susceptibility loci for colorectal cancer. <i>Nature Genetics</i> , 2008, 40, 1426-1435.	9.4	498
9	Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. <i>Human Molecular Genetics</i> , 2008, 17, 3720-3727.	1.4	61
10	Association of a Common AKAP9 Variant With Breast Cancer Risk: A Collaborative Analysis. <i>Journal of the National Cancer Institute</i> , 2008, 100, 437-442.	3.0	44
11	Functional Polymorphisms in Folate Metabolism Genes Influence the Risk of Meningioma and Glioma. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2008, 17, 1195-1202.	1.1	48
12	Counting potentially functional variants in BRCA1, BRCA2 and ATM predicts breast cancer susceptibility. <i>Human Molecular Genetics</i> , 2007, 16, 1051-1057.	1.4	109
13	A genome-wide association study shows that common alleles of SMAD7 influence colorectal cancer risk. <i>Nature Genetics</i> , 2007, 39, 1315-1317.	9.4	463
14	A genome-wide association scan of tag SNPs identifies a susceptibility variant for colorectal cancer at 8q24.21. <i>Nature Genetics</i> , 2007, 39, 984-988.	9.4	754