Emily Webb

List of Publications by Year
in descending order

14 papers	3,753 citations	687363 13 h-index	1058476 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. Genes Chromosomes and Cancer, 2010, 49, 539-548.	2.8	17
2	The colorectal cancer risk at 18q21 is caused by a novel variant altering *SMAD7* expression. Genome Research, 2009, 19, 987-993.	5.5	85
3	A genome-wide scan of 10â€‰000 gene-centric variants and colorectal cancer risk. European Journal of Human Genetics, 2009, 17, 1507-1514.	2.8	12
4	A genome-wide association study identifies colorectal cancer susceptibility loci on chromosomes 10p14 and 8q23.3. Nature Genetics, 2008, 40, 623-630.	21.4	514
5	Genome-wide association scan identifies a colorectal cancer susceptibility locus on 11q23 and replicates risk loci at 8q24 and 18q21. Nature Genetics, 2008, 40, 631-637.	21.4	542
6	Common genetic variants at the CRAC1 (HMPS) locus on chromosome 15q13.3 influence colorectal cancer risk. Nature Genetics, 2008, 40, 26-28.	21.4	277
7	A genome-wide association study identifies six susceptibility loci for chronic lymphocytic leukemia. Nature Genetics, 2008, 40, 1204-1210.	21.4	329
8	Meta-analysis of genome-wide association data identifies four new susceptibility loci for colorectal cancer. Nature Genetics, 2008, 40, 1426-1435.	21.4	498
9	Refinement of the basis and impact of common 11q23.1 variation to the risk of developing colorectal cancer. Human Molecular Genetics, 2008, 17, 3720-3727.	2.9	61
10	Association of a Common AKAP9 Variant With Breast Cancer Risk: A Collaborative Analysis. Journal of the National Cancer Institute, 2008, 100, 437-442.	6.3	44
11	Functional Polymorphisms in Folate Metabolism Genes Influence the Risk of Meningioma and Glioma. Cancer Epidemiology Biomarkers and Prevention, 2008, 17, 1195-1202.	2.5	48
12	Counting potentially functional variants in BRCA1, BRCA2 and ATM predicts breast cancer susceptibility. Human Molecular Genetics, 2007, 16, 1051-1057.	2.9	109
13	A genome-wide association study shows that common alleles of SMAD7 influence colorectal cancer risk. Nature Genetics, 2007, 39, 1315-1317.	21.4	463
14	A genome-wide association scan of tag SNPs identifies a susceptibility variant for colorectal cancer at 8q24.21. Nature Genetics, 2007, 39, 984-988.	21.4	754