Recurrent Coding Sequence Variation Explains Only A Small Fraction of the Genetic Architecture of Colorectal Cancer

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#	Paper	IF	Citations
23	Correspondence: SEMA4A variation and risk of colorectal cancer. *Nature Communications*, **2016**, 7, 10611	17.4	6
22	Somatic POLE proofreading domain mutation, immune response, and prognosis in colorectal cancer: a retrospective, pooled biomarker study. *The Lancet Gastroenterology and Hepatology*, **2016**, 1, 207-216	18.8	160
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19	Validation of Recently Proposed Colorectal Cancer Susceptibility Gene Variants in an Analysis of Families and Patients-a Systematic Review. *Gastroenterology*, **2017**, 152, 75-77.e4	13.3	62
18	Coding variants in NOD-like receptors: An association study on risk and survival of colorectal cancer. *PLoS ONE*, **2018**, 13, e0199350	3.7	5
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16	Genome-wide scan of the effect of common nsSNPs on colorectal cancer survival outcome. *British Journal of Cancer*, **2018**, 119, 988-993	8.7	4
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