

Lori S Tillmans

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Cytotoxic T Cells and Granzyme B Associated with Improved Colorectal Cancer Survival in a Prospective Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2017, 26, 622-631.	2.5	68
2	High-throughput screening of prostate cancer risk loci by single nucleotide polymorphisms sequencing. <i>Nature Communications</i> , 2018, 9, 2022.	12.8	66
3	Tumor eosinophil infiltration and improved survival of colorectal cancer patients: Iowa Women's Health Study. <i>Modern Pathology</i> , 2016, 29, 516-527.	5.5	65
4	Prostate cancer risk locus at 8q24 as a regulatory hub by physical interactions with multiple genomic loci across the genome. <i>Human Molecular Genetics</i> , 2015, 24, 154-166.	2.9	53
5	Incorporating Functional Annotations for Fine-Mapping Causal Variants in a Bayesian Framework Using Summary Statistics. <i>Genetics</i> , 2016, 204, 933-958.	2.9	51
6	Chromatin interactions and candidate genes at ten prostate cancer risk loci. <i>Scientific Reports</i> , 2016, 6, 23202.	3.3	36
7	Biallelic BRCA2 Mutations Shape the Somatic Mutational Landscape of Aggressive Prostate Tumors. <i>American Journal of Human Genetics</i> , 2016, 98, 818-829.	6.2	34
8	Associations between Cigarette Smoking, Hormone Therapy, and Folate Intake with Incident Colorectal Cancer by TP53 Protein Expression Level in a Population-Based Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2014, 23, 350-355.	2.5	11
9	Associations between Environmental Exposures and Incident Colorectal Cancer by ESR2 Protein Expression Level in a Population-Based Cohort of Older Women. <i>Cancer Epidemiology Biomarkers and Prevention</i> , 2015, 24, 713-719.	2.5	10
10	Single-Nucleotide Polymorphisms Sequencing Identifies Candidate Functional Variants at Prostate Cancer Risk Loci. <i>Genes</i> , 2019, 10, 547.	2.4	7
11	An expanded variant list and assembly annotation identifies multiple novel coding and noncoding genes for prostate cancer risk using a normal prostate tissue eQTL data set. <i>PLoS ONE</i> , 2019, 14, e0214588.	2.5	5
12	A microRNA Transcriptome-wide Association Study of Prostate Cancer Risk. <i>Frontiers in Genetics</i> , 2022, 13, 836841.	2.3	3
13	Associations between tissue-based CD3+ T lymphocyte count and colorectal cancer survival in a prospective cohort of older women. <i>Molecular Carcinogenesis</i> , 2021, 60, 15-24.	2.7	1