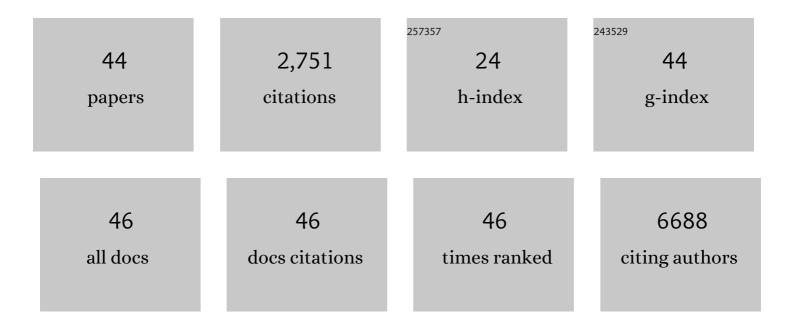
## Maarit Tiirikainen

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Beta-catenin activation and immunotherapy resistance in hepatocellular carcinoma: mechanisms and biomarkers. Hepatoma Research, 2021, 2021, .	0.6	14
2	Genome-wide epigenetic analyses in Japanese immigrant plantation workers with Parkinson's disease and exposure to organochlorines reveal possible involvement of glial genes and pathways involved in neurotoxicity. BMC Neuroscience, 2020, 21, 31.	0.8	16
3	Population-specific reference panels are crucial for genetic analyses: an example of the CREBRF locus in Native Hawaiians. Human Molecular Genetics, 2020, 29, 2275-2284.	1.4	27
4	Diagnostic Power of DNA Methylation Classifiers for Early Detection of Cancer. Trends in Cancer, 2020, 6, 78-81.	3.8	58
5	Transcriptomics Associates Molecular Features with 18F-Fluorocholine PET/CT Imaging Phenotype and Its Potential Relationship to Survival in Hepatocellular Carcinoma. Cancer Research, 2019, 79, 1696-1704.	0.4	13
6	Influence of UGT2B10 Genotype on Urinary Excretion of 4-(Methylnitrosamino)-1-(3-pyridyl)-1-butanol- <i>N-</i> glucuronide by African American Smokers. Chemical Research in Toxicology, 2018, 31, 168-175.	1.7	4
7	Clinical and molecular sub-classification of hepatocellular carcinoma relative to alpha-fetoprotein level in an Asia-Pacific island cohort. Hepatoma Research, 2018, 4, 1.	0.6	14
8	Methylation of imprinted <i>IGF2</i> regions is associated with total, visceral, and hepatic adiposity in postmenopausal women. Epigenetics, 2018, 13, 858-865.	1.3	6
9	Association of internal smoking dose with blood DNA methylation in three racial/ethnic populations. Clinical Epigenetics, 2018, 10, 110.	1.8	34
10	Genetic Determinants of 1,3-Butadiene Metabolism and Detoxification in Three Populations of Smokers with Different Risks of Lung Cancer. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 1034-1042.	1.1	22
11	Comparison of genome-scale DNA methylation profiles in hepatocellular carcinoma by viral status. Epigenetics, 2016, 11, 464-474.	1.3	15
12	Pan-Cancer Analyses Reveal Long Intergenic Non-Coding RNAs Relevant to Tumor Diagnosis, Subtyping and Prognosis. EBioMedicine, 2016, 7, 62-72.	2.7	33
13	Genetic determinants of CYP2A6 activity across racial/ethnic groups with different risks of lung cancer and effect on their smoking intensity. Carcinogenesis, 2016, 37, 269-279.	1.3	48
14	Metabolic characteristics distinguishing intrahepatic cholangiocarcinoma: a negative pilot study of (18)F-fluorocholine PET/CT clarified by transcriptomic analysis. American Journal of Nuclear Medicine and Molecular Imaging, 2016, 6, 73-83.	1.0	5
15	High Incidence of Somatic BAP1 Alterations in Sporadic Malignant Mesothelioma. Journal of Thoracic Oncology, 2015, 10, 565-576.	0.5	282
16	Intraindividual Variation and Short-term Temporal Trend in DNA Methylation of Human Blood. Cancer Epidemiology Biomarkers and Prevention, 2015, 24, 490-497.	1.1	27
17	Genome-scale hypomethylation in the cord blood DNAs associated with early onset preeclampsia. Clinical Epigenetics, 2015, 7, 21.	1.8	41
18	Pleiotropy of Cancer Susceptibility Variants on the Risk of Non-Hodgkin Lymphoma: The PAGE Consortium. PLoS ONE, 2014, 9, e89791.	1.1	16

MAARIT TIIRIKAINEN

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19	1,3-Butadiene Exposure and Metabolism among Japanese American, Native Hawaiian, and White Smokers. Cancer Epidemiology Biomarkers and Prevention, 2014, 23, 2240-2249.	1.1	22
20	Genome-wide hypermethylation coupled with promoter hypomethylation in the chorioamniotic membranes of early onset pre-eclampsia. Molecular Human Reproduction, 2014, 20, 885-904.	1.3	54
21	<i>CCAT2</i> , a novel noncoding RNA mapping to 8q24, underlies metastatic progression and chromosomal instability in colon cancer. Genome Research, 2013, 23, 1446-1461.	2.4	526
22	Genetic associations of relaxin: preterm birth and premature rupture of fetal membranes. American Journal of Obstetrics and Gynecology, 2013, 209, 258.e1-258.e8.	0.7	19
23	Continuous Exposure to Chrysotile Asbestos Can Cause Transformation of Human Mesothelial Cells via HMCB1 and TNF-α Signaling. American Journal of Pathology, 2013, 183, 1654-1666.	1.9	88
24	Integrated analysis of genomeâ€wide copy number alterations and gene expression in microsatellite stable, CpG island methylator phenotypeâ€negative colon cancer. Genes Chromosomes and Cancer, 2013, 52, 450-466.	1.5	51
25	Generalization and Dilution of Association Results from European GWAS in Populations of Non-European Ancestry: The PAGE Study. PLoS Biology, 2013, 11, e1001661.	2.6	235
26	Elucidating the Landscape of Aberrant DNA Methylation in Hepatocellular Carcinoma. PLoS ONE, 2013, 8, e55761.	1.1	101
27	cis-Expression QTL Analysis of Established Colorectal Cancer Risk Variants in Colon Tumors and Adjacent Normal Tissue. PLoS ONE, 2012, 7, e30477.	1.1	55
28	Susceptibility variants for obesity and colorectal cancer risk: The multiethnic cohort and PAGE studies. International Journal of Cancer, 2012, 131, E1038-43.	2.3	31
29	Measurement of Circulating Cellâ€Free DNA in Relation to 18Fâ€Fluorocholine PET/CT Imaging in Chemotherapyâ€Treated Advanced Prostate Cancer. Clinical and Translational Science, 2012, 5, 65-70.	1.5	49
30	Type 2 diabetes risk variants and colorectal cancer risk: the Multiethnic Cohort and PAGE studies. Gut, 2011, 60, 1703-1711.	6.1	43
31	ZEB1 limits adenoviral infectability by transcriptionally repressing the Coxsackie virus and Adenovirus Receptor. Molecular Cancer, 2011, 10, 91.	7.9	19
32	Persistent Inflammatory Pathways Associated with Early Onset Myocardial Infarction in a Medicated Multiethnic Hawaiian Cohort. Biochemistry Insights, 2011, 4, BCI.S6976.	3.3	10
33	Urinary Isothiocyanates; Glutathione <i>S</i> -Transferase <i>M1, T1</i> , and <i>P1</i> Polymorphisms; and Risk of Colorectal Cancer: The Multiethnic Cohort Study. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 314-320.	1.1	41
34	Plasma Levels of B Vitamins and Colorectal Cancer Risk: The Multiethnic Cohort Study. Cancer Epidemiology Biomarkers and Prevention, 2009, 18, 2195-2201.	1.1	72
35	Smokers with the CHRNA Lung Cancer–Associated Variants Are Exposed to Higher Levels of Nicotine Equivalents and a Carcinogenic Tobacco-Specific Nitrosamine. Cancer Research, 2008, 68, 9137-9140.	0.4	186
36	Variants on 9p24 and 8q24 Are Associated with Risk of Colorectal Cancer: Results from the Colon Cancer Family Registry. Cancer Research, 2007, 67, 11128-11132.	0.4	87

Maarit Tiirikainen

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37	Transforming Growth Factor-β Receptor Inhibition Enhances Adenoviral Infectability of Carcinoma Cells via Up-Regulation of Coxsackie and Adenovirus Receptor in Conjunction with Reversal of Epithelial-Mesenchymal Transition. Cancer Research, 2006, 66, 1648-1657.	0.4	68
38	Genomic alterations in human mesothelioma including high resolution mapping of common regions of DNA loss in chromosome arm 6q. Anticancer Research, 2003, 23, 2281-9.	0.5	22
39	Integrated genomic and epigenomic analyses pinpoint biallelic gene inactivation in tumors. Nature Genetics, 2002, 32, 453-458.	9.4	172
40	INITIAL P-GLYCOPROTEIN EXPRESSION IN CHILDHOOD ACUTE LYMPHOBLASTIC LEUKEMIA: NO EVIDENCE OF PROGNOSTIC IMPACT IN FOLLOW-UP. Pediatric Hematology and Oncology, 2001, 18, 27-36.	0.3	21
41	DNA Copy Number Alterations in HIV-Positive and HIV-Negative Patients With Diffuse Large-Cell Lymphomas. Journal of Acquired Immune Deficiency Syndromes (1999), 2001, 27, 272-276.	0.9	8
42	Multiple Drug Resistance Mediated by P-Glycoprotein is not a Major Factor in a Slow Response to Therapy in Childhood All. Pediatric Hematology and Oncology, 1998, 15, 11-21.	0.3	23
43	Evaluation of red blood cell lysing solutions for the detection of intracellular antigens by flow cytometry. Cytometry, 1995, 20, 341-348.	1.8	50
44	Flow Cytometric Analysis of Terminal Deoxynucleotidyl Transferase: <i>A Simplified Method</i> . American Journal of Clinical Pathology, 1993, 99, 298-303.	0.4	23