Lee E Moore

List of Publications by Year
in descending order

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45 papers	2,427 citations	201674 27 h-index	223800 46 g-index
46 all docs	46 docs citations	46 times ranked	5412 citing authors

#	Article	IF	Citations
1	Genome-wide association study of renal cell carcinoma identifies two susceptibility loci on 2p21 and 11q13.3. Nature Genetics, 2011, 43, 60-65.	21.4	220
2	Genomic DNA hypomethylation as a biomarker for bladder cancer susceptibility in the Spanish Bladder Cancer Study: a caseâ€“control study. Lancet Oncology, The, 2008, 9, 359-366.	10.7	211
3	Von Hippel-Lindau (VHL) Inactivation in Sporadic Clear Cell Renal Cancer: Associations with Germline VHL Polymorphisms and Etiologic Risk Factors. PLoS Genetics, 2011, 7, e1002312.	3.5	168
4	Analysis of Heritability and Shared Heritability Based on Genome-Wide Association Studies for Thirteen Cancer Types. Journal of the National Cancer Institute, 2015, 107, djv279.	6.3	152
5	Lifestyle Factors, Exposures, Genetic Susceptibility, and Renal Cell Cancer Risk: A Review. Cancer Investigation, 2005, 23, 240-255.	1.3	110
6	Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724.	12.8	106
7	Elevated Bladder Cancer in Northern New England: The Role of Drinking Water and Arsenic. Journal of the National Cancer Institute, 2016, 108, .	6.3	102
8	Concurrent Alterations in TERT, KDM6A, and the BRCA Pathway in Bladder Cancer. Clinical Cancer Research, 2014, 20, 4935-4948.	7.0	101
9	Occupational Trichloroethylene Exposure and Renal Carcinoma Risk: Evidence of Genetic Susceptibility by Reductive Metabolism Gene Variants. Cancer Research, 2010, 70, 6527-6536.	0.9	97
10	Obesity and cancer: Mendelian randomization approach utilizing the FTO genotype. International Journal of Epidemiology, 2009, 38, 971-975.	1.9	96
11	Female chromosome X mosaicism is age-related and preferentially affects the inactivated X chromosome. Nature Communications, 2016, 7, 11843.	12.8	86
12	Evaluation of Apolipoprotein A1 and Posttranslationally Modified Forms of Transthyretin as Biomarkers for Ovarian Cancer Detection in an Independent Study Population. Cancer Epidemiology Biomarkers and Prevention, 2006, 15, 1641-1646.	2.5	85
13	A genome-wide association study identifies a novel susceptibility locus for renal cell carcinoma on 12p11.23. Human Molecular Genetics, 2012, 21, 456-462.	2.9	81
14	Proteomic biomarkers in combination with CA 125 for detection of epithelial ovarian cancer using prediagnostic serum samples from the Prostate, Lung, Colorectal, and Ovarian (PLCO) Cancer Screening Trial. Cancer, 2012, 118, 91-100.	4.1	77
15	Polymorphisms in one-carbon metabolism and trans-sulfuration pathway genes and susceptibility to bladder cancer. International Journal of Cancer, 2007, 120, 2452-2458.	5.1	60
16	The influence of obesity-related factors in the etiology of renal cell carcinomaâ€”A mendelian randomization study. PLoS Medicine, 2019, 16, e1002724.	8.4	59
17	High pesticide exposure events and <scp>DNA</scp> methylation among pesticide applicators in the agricultural health study. Environmental and Molecular Mutagenesis, 2017, 58, 19-29.	2.2	48
18	Epidemiologic Considerations to Assess Altered DNA Methylation from Environmental Exposures in Cancer. Annals of the New York Academy of Sciences, 2003, 983, 181-196.	3.8	43

#	Article	IF	Citations
19	A case-control study of occupational exposure to metalworking fluids and bladder cancer risk among men. Occupational and Environmental Medicine, 2014, 71, 667-674.	2.8	43
20	Genetic Variants Related to Longer Telomere Length are Associated with Increased Risk of Renal Cell Carcinoma. European Urology, 2017, 72, 747-754.	1.9	39
21	The chromosome 2p21 region harbors a complex genetic architecture for association with risk for renal cell carcinoma. Human Molecular Genetics, 2012, 21, 1190-1200.	2.9	37
22	LINE1methylation levels associated with increased bladder cancer risk in pre-diagnostic blood DNA among US (PLCO) and European (ATBC) cohort study participants. Epigenetics, 2014, 9, 404-415.	2.7	35
23	Modification of Occupational Exposures on Bladder Cancer Risk by Common Genetic Polymorphisms. Journal of the National Cancer Institute, 2015, 107, djv223.	6.3	34
24	Folate metabolism genes, vegetable intake and renal cancer risk in central Europe. International Journal of Cancer, 2008, 122, 1710-1715.	5.1	33
25	Apolipoprotein E/C1 Locus Variants Modify Renal Cell Carcinoma Risk. Cancer Research, 2009, 69, 8001-8008.	0.9	31
26	Prospective study of DNA methylation at LINEâ€1 and Alu in peripheral blood and the risk of prostate cancer. Prostate, 2015, 75, 1718-1725.	2.3	30
27	Parental occupational exposures and Ewing's sarcoma. International Journal of Cancer, 2005, 114, 472-478.	5.1	27
28	Sex specific associations in genome wide association analysis of renal cell carcinoma. European Journal of Human Genetics, 2019, 27, 1589-1598.	2.8	27
29	Potential effect modifiers of the arsenicâ€“bladder cancer risk relationship. International Journal of Cancer, 2018, 143, 2640-2646.	5.1	25
30	Diesel exhaust and bladder cancer risk by pathologic stage and grade subtypes. Environment International, 2020, 135, 105346.	10.0	25
31	The 19q12 Bladder Cancer GWAS Signal: Association with Cyclin E Function and Aggressive Disease. Cancer Research, 2014, 74, 5808-5818.	0.9	24
32	Smoking status, usual adult occupation, and risk of recurrent urothelial bladder carcinoma: data from The Cancer Genome Atlas (TCGA) Project. Cancer Causes and Control, 2016, 27, 1429-1435.	1.8	18
33	Pesticide use and LINE-1 methylation among male private pesticide applicators in the Agricultural Health Study. Environmental Epigenetics, 2017, 3, dvx005.	1.8	16
34	Prospective study of DNA methylation at chromosome 8q24 in peripheral blood and prostate cancer risk. British Journal of Cancer, 2017, 116, 1470-1479.	6.4	15
35	Analgesic use and risk of renal cell carcinoma: A case-control, cohort and meta-analytic assessment. International Journal of Cancer, 2016, 139, 584-592.	5.1	11
36	Targeted Deep Sequencing of Bladder Tumors Reveals Novel Associations between Cancer Gene Mutations and Mutational Signatures with Major Risk Factors. Clinical Cancer Research, 2021, 27, 3725-3733.	7.0	11

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37	Editorâ€™s Highlight: High-Throughput Functional Genomics Identifies Modulators of TCE Metabolite Genotoxicity and Candidate Susceptibility Genes. Toxicological Sciences, 2017, 160, 111-120.	3.1	10
38	Use of OMIC technologies to study arsenic exposure in human populations. Environmental and Molecular Mutagenesis, 2013, 54, 589-595.	2.2	9
39	A caseâ€“control study of occupational sunlight exposure and renal cancer risk. International Journal of Cancer, 2016, 138, 1626-1633.	5.1	8
40	Differences in risk factors for molecular subtypes of clear cell renal cell carcinoma. International Journal of Cancer, 2021, 149, 1448-1454.	5.1	5
41	Analysis of the Distribution and Temporal Trends of Grade and Stage in Urothelial Bladder Cancer in Northern New England from 1994 to 2004. ISRN Pathology, 2012, 2012, 1-7.	0.4	4
42	Differences in Tumor VHL Mutation and Hypoxia-inducible Factor 2Î± Expression Between African American and White Patients with Clear Cell Renal Cell Carcinoma. European Urology, 2019, 75, 882-884.	1.9	3
43	Coffee intake and trace element blood concentrations in association with renal cell cancer among smokers. Cancer Causes and Control, 2022, 33, 91-99.	1.8	2
44	0084â€…A Case-Control Study of Occupational Exposure to Metalworking Fluids and Bladder Cancer Risk among Men. Occupational and Environmental Medicine, 2014, 71, A71.1-A71.	2.8	1
45	RE: Elevated Bladder Cancer in Northern New England: The Role of Drinking Water and Arsenic. Journal of the National Cancer Institute, 2018, 110, 1273-1274.	6.3	1