# James D Mckay

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/3251254/publications.pdf

Version: 2024-02-01

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|---|---|---|---|
| 40<br>papers | 2,600<br>citations | 279798<br>23<br>h-index | 289244<br>40<br>g-index |
| 42<br>all docs | 42<br>docs citations | 42<br>times ranked | 6329<br>citing authors |

| # | Article | IF | Citations |
|---|---|---|---|
| 1 | Development of a custom next-generation sequencing panel for the determination of bladder cancer risk in a Tunisian cohort. Molecular Biology Reports, 2022, 49, 1233-1258. | 2.3 | 4 |
| 2 | Genome-wide interaction analysis identified low-frequency variants with sex disparity in lung cancer risk. Human Molecular Genetics, 2022, 31, 2831-2843. | 2.9 | 4 |
| 3 | Genetic Analysis of Lung Cancer and the Germline Impact on Somatic Mutation Burden. Journal of the National Cancer Institute, 2022, 114, 1159-1166. | 6.3 | 8 |
| 4 | Circulating Isovalerylcarnitine and Lung Cancer Risk: Evidence from Mendelian Randomization and Prediagnostic Blood Measurements. Cancer Epidemiology Biomarkers and Prevention, 2022, 31, 1966-1974. | 2.5 | 4 |
| 5 | Association between anthropometry and lifestyle factors and risk of Bâ€cell lymphoma: An exposomeâ€wide analysis. International Journal of Cancer, 2021, 148, 2115-2128. | 5.1 | 9 |
| 6 | Sexual dimorphism in cancer: insights from transcriptional signatures in kidney tissue and renal cell carcinoma. Human Molecular Genetics, 2021, 30, 343-355. | 2.9 | 14 |
| 7 | Sequencing at lymphoid neoplasm susceptibility loci maps six myeloma risk genes. Human Molecular Genetics, 2021, 30, 1142-1153. | 2.9 | 2 |
| 8 | The Shared Genetic Architectures Between Lung Cancer and Multiple Polygenic Phenotypes in Genome-Wide Association Studies. Cancer Epidemiology Biomarkers and Prevention, 2021, 30, 1156-1164. | 2.5 | 13 |
| 9 | Genome-wide association meta-analysis identifies pleiotropic risk loci for aerodigestive squamous cell cancers. PLoS Genetics, 2021, 17, e1009254. | 3.5 | 19 |
| 10 | The PI3K/mTOR Pathway Is Targeted by Rare Germline Variants in Patients with Both Melanoma and Renal Cell Carcinoma. Cancers, 2021, 13, 2243. | 3.7 | 2 |
| 11 | The shared genetic architecture between epidemiological and behavioral traits with lung cancer. Scientific Reports, 2021, 11, 17559. | 3.3 | 10 |
| 12 | Germline determinants of humoral immune response to HPV-16 protect against oropharyngeal cancer. Nature Communications, 2021, 12, 5945. | 12.8 | 10 |
| 13 | Mediating effect of soluble B-cell activation immune markers on the association between anthropometric and lifestyle factors and lymphoma development. Scientific Reports, 2020, 10, 13814. | 3.3 | 4 |
| 14 | A molecular map of lung neuroendocrine neoplasms. GigaScience, 2020, 9, . | 6.4 | 17 |
| 15 | Needlestack: an ultra-sensitive variant caller for multi-sample next generation sequencing data. NAR Genomics and Bioinformatics, 2020, 2, lqaa021. | 3.2 | 5 |
| 16 | Urinary TERT promoter mutations are detectable up to 10 years prior to clinical diagnosis of bladder cancer: Evidence from the Golestan Cohort Study. EBioMedicine, 2020, 53, 102643. | 6.1 | 51 |
| 17 | Exome sequencing identifies germline variants in DIS3 in familial multiple myeloma. Leukemia, 2019, 33, 2324-2330. | 7.2 | 33 |
| 18 | Redefining malignant pleural mesothelioma types as a continuum uncovers immune-vascular interactions. EBioMedicine, 2019, 48, 191-202. | 6.1 | 55 |

| # | Article | IF | Citations |
|---|---|---|---|
| 19 | 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 |
| 20 | Mendelian Randomization and mediation analysis of leukocyte telomere length and risk of lung and head and neck cancers. International Journal of Epidemiology, 2019, 48, 751-766. | 1.9 | 32 |
| 21 | Integrative genomic profiling of large-cell neuroendocrine carcinomas reveals distinct subtypes of high-grade neuroendocrine lung tumors. Nature Communications, 2018, 9, 1048. | 12.8 | 254 |
| 22 | Novel pedigree analysis implicates DNA repair and chromatin remodeling in multiple myeloma risk. PLoS Genetics, 2018, 14, e1007111. | 3.5 | 30 |
| 23 | Genome-wide association study identifies multiple risk loci for renal cell carcinoma. Nature Communications, 2017, 8, 15724. | 12.8 | 106 |
| 24 | Large-scale association analysis identifies new lung cancer susceptibility loci and heterogeneity in genetic susceptibility across histological subtypes. Nature Genetics, 2017, 49, 1126-1132. | 21.4 | 472 |
| 25 | Obesity, metabolic factors and risk of different histological types of lung cancer: A Mendelian randomization study. PLoS ONE, 2017, 12, e0177875. | 2.5 | 79 |
| 26 | Identification of Circulating Tumor DNA for the Early Detection of Small-cell Lung Cancer. EBioMedicine, 2016, 10, 117-123. | 6.1 | 153 |
| 27 | The causal relevance of body mass index in different histological types of lung cancer: A Mendelian randomization study. Scientific Reports, 2016, 6, 31121. | 3.3 | 27 |
| 28 | KRASmutations in blood circulating cell-free DNA: a pancreatic cancer case-control. Oncotarget, 2016, 7, 78827-78840. | 1.8 | 70 |
| 29 | A genome-wide association study of marginal zone lymphoma shows association to the HLA region. Nature Communications, 2015, 6, 5751. | 12.8 | 58 |
| 30 | A Rare Truncating BRCA2 Variant and Genetic Susceptibility to Upper Aerodigestive Tract Cancer. Journal of the National Cancer Institute, 2015, 107, . | 6.3 | 33 |
| 31 | Identification of lung cancer histology-specific variants applying Bayesian framework variant prioritization approaches within the TRICL and ILCCO consortia. Carcinogenesis, 2015, 36, 1314-1326. | 2.8 | 15 |
| 32 | Common Variation at 1q24.1 (ALDH9A1) Is a Potential Risk Factor for Renal Cancer. PLoS ONE, 2015, 10, e0122589. | 2.5 | 19 |
| 33 | Genome-wide association study identifies multiple susceptibility loci for diffuse large B cell lymphoma. Nature Genetics, 2014, 46, 1233-1238. | 21.4 | 147 |
| 34 | Genome-wide Association Study Identifies Five Susceptibility Loci for Follicular Lymphoma outside the HLA Region. American Journal of Human Genetics, 2014, 95, 462-471. | 6.2 | 96 |
| 35 | Common variation at 2q22.3 (ZEB2) influences the risk of renal cancer. Human Molecular Genetics, 2013, 22, 825-831. | 2.9 | 54 |
| 36 | Influence of common genetic variation on lung cancer risk: meta-analysis of 14 900 cases and 29 485 controls. Human Molecular Genetics, 2012, 21, 4980-4995. | 2.9 | 196 |

| # | Article | IF | Citations |
|---|---|---|---|
| 37 | 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 |
| 38 | 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 |
| 39 | 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 |
| 40 | Obesity and cancer: Mendelian randomization approach utilizing the FTO genotype. International Journal of Epidemiology, 2009, 38, 971-975. | 1.9 | 96 |