## Liam F Spurr

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/701266/publications.pdf

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|          |                | 759233       | 794594         |
|----------|----------------|--------------|----------------|
| 18       | 1,130          | 12           | 19             |
| papers   | citations      | h-index      | g-index        |
|          |                |              |                |
|          |                |              |                |
|          |                |              |                |
| 24       | 24             | 24           | 2048           |
| all docs | docs citations | times ranked | citing authors |
|          |                |              |                |

| #  | Article   | IF   | Citations |
|----|---|------|-----------|
| 1  | Quantification of aneuploidy in targeted sequencing data using ASCETS. Bioinformatics, 2021, 37, 2461-2463.   | 4.1  | 21        |
| 2  | Genomic Characterization of <i>de novo</i> Metastatic Breast Cancer. Clinical Cancer Research, 2021, 27, 1105-1118.   | 7.0  | 24        |
| 3  | Molecular Characterization and Therapeutic Targeting of Colorectal Cancers Harboring Receptor<br>Tyrosine Kinase Fusions. Clinical Cancer Research, 2021, 27, 1695-1705.  | 7.0  | 19        |
| 4  | SMARCA4 and Other SWItch/Sucrose NonFermentable Family Genomic Alterations in NSCLC: Clinicopathologic Characteristics and Outcomes to Immune Checkpoint Inhibition. Journal of Thoracic Oncology, 2021, 16, 1176-1187. | 1.1  | 49        |
| 5  | scReQTL: an approach to correlate SNVs to gene expression from individual scRNA-seq datasets. BMC Genomics, 2021, 22, 40.   | 2.8  | 11        |
| 6  | CDKN2A loss-of-function predicts immunotherapy resistance in non-small cell lung cancer. Scientific Reports, 2021, 11, 20059.   | 3.3  | 53        |
| 7  | Microenvironment drives cell state, plasticity, and drug response in pancreatic cancer. Cell, 2021, 184, 6119-6137.e26.   | 28.9 | 201       |
| 8  | ReQTL: identifying correlations between expressed SNVs and gene expression using RNA-sequencing data. Bioinformatics, 2020, 36, 1351-1359.  | 4.1  | 13        |
| 9  | GeTallele: A Method for Analysis of DNA and RNA Allele Frequency Distributions. Frontiers in Bioengineering and Biotechnology, 2020, 8, 1021.   | 4.1  | 2         |
| 10 | Synthetic Lethal Interaction between the ESCRT Paralog Enzymes VPS4A and VPS4B in Cancers Harboring Loss of Chromosome 18q or 16q. Cell Reports, 2020, 33, 108493.  | 6.4  | 28        |
| 11 | Impact of DNA Damage Response and Repair (DDR) Gene Mutations on Efficacy of PD-(L)1 Immune<br>Checkpoint Inhibition in Non–Small Cell Lung Cancer. Clinical Cancer Research, 2020, 26, 4135-4142.                      | 7.0  | 95        |
| 12 | Molecular Mechanisms of Acquired Resistance to MET Tyrosine Kinase Inhibitors in Patients with MET Exon 14–Mutant NSCLC. Clinical Cancer Research, 2020, 26, 2615-2625.   | 7.0  | 129       |
| 13 | Transcriptional dysregulation in developing trigeminal sensory neurons in the LgDel mouse model of DiGeorge 22q11.2 deletion syndrome. Human Molecular Genetics, 2020, 29, 1002-1017.                                   | 2.9  | 8         |
| 14 | Estimating the Allele-Specific Expression of SNVs From $10\tilde{A}$ — Genomics Single-Cell RNA-Sequencing Data. Genes, 2020, $11$ , 240.   | 2.4  | 14        |
| 15 | Mammalian SWI/SNF Complex Genomic Alterations and Immune Checkpoint Blockade in Solid Tumors.<br>Cancer Immunology Research, 2020, 8, 1075-1084.  | 3.4  | 47        |
| 16 | Mechanisms and therapeutic implications of hypermutation in gliomas. Nature, 2020, 580, 517-523.  | 27.8 | 374       |
| 17 | Systematic pan-cancer analysis of somatic allele frequency. Scientific Reports, 2018, 8, 7735.  | 3.3  | 21        |
| 18 | Overexpressed somatic alleles are enriched in functional elements in Breast Cancer. Scientific Reports, 2017, 7, 8287.  | 3.3  | 3         |