

Nathan Lawlor

List of Publications by Year in descending order

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Version: 2024-02-01

16
papers

1,162
citations

840119

11
h-index

940134

16
g-index

22
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22
docs citations

22
times ranked

2209
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Single-cell transcriptomes identify human islet cell signatures and reveal cell-type-specific expression changes in type 2 diabetes. <i>Genome Research</i> , 2017, 27, 208-222. | 2.4 | 440 |
| 2 | The human milk oligosaccharide 2-fucosyllactose modulates CD14 expression in human enterocytes, thereby attenuating LPS-induced inflammation. <i>Gut</i> , 2016, 65, 33-46. | 6.1 | 217 |
| 3 | Human Milk Components Modulate Toll-Like Receptor-Mediated Inflammation. <i>Advances in Nutrition</i> , 2016, 7, 102-111. | 2.9 | 114 |
| 4 | Multiomic Profiling Identifies cis-Regulatory Networks Underlying Human Pancreatic β^2 Cell Identity and Function. <i>Cell Reports</i> , 2019, 26, 788-801.e6. | 2.9 | 68 |
| 5 | Genomics of Islet (Dys)function and Type 2 Diabetes. <i>Trends in Genetics</i> , 2017, 33, 244-255. | 2.9 | 55 |
| 6 | A Common Type 2 Diabetes Risk Variant Potentiates Activity of an Evolutionarily Conserved Islet Stretch Enhancer and Increases C2CD4A and C2CD4B Expression. <i>American Journal of Human Genetics</i> , 2018, 102, 620-635. | 2.6 | 47 |
| 7 | Type 2 Diabetes-Associated Genetic Variants Regulate Chromatin Accessibility in Human Islets. <i>Diabetes</i> , 2018, 67, 2466-2477. | 0.3 | 44 |
| 8 | Alpha TC1 and Beta-TC-6 genomic profiling uncovers both shared and distinct transcriptional regulatory features with their primary islet counterparts. <i>Scientific Reports</i> , 2017, 7, 11959. | 1.6 | 41 |
| 9 | AMULET: a novel read count-based method for effective multiplet detection from single nucleus ATAC-seq data. <i>Genome Biology</i> , 2021, 22, 252. | 3.8 | 36 |
| 10 | Single Cell Analysis of Blood Mononuclear Cells Stimulated Through Either LPS or Anti-CD3 and Anti-CD28. <i>Frontiers in Immunology</i> , 2021, 12, 636720. | 2.2 | 32 |
| 11 | multiClust: An R-package for Identifying Biologically Relevant Clusters in Cancer Transcriptome Profiles. <i>Cancer Informatics</i> , 2016, 15, CIN.S38000. | 0.9 | 14 |
| 12 | (Epi)genomic heterogeneity of pancreatic islet function and failure in type 2 diabetes. <i>Molecular Metabolism</i> , 2019, 27, S15-S24. | 3.0 | 12 |
| 13 | Tet2 Controls the Responses of β^2 cells to Inflammation in Autoimmune Diabetes. <i>Nature Communications</i> , 2021, 12, 5074. | 5.8 | 11 |
| 14 | BiFET: sequencing bias-free transcription factor footprint enrichment test. <i>Nucleic Acids Research</i> , 2019, 47, e11-e11. | 6.5 | 9 |
| 15 | Detection of correlated hidden factors from single cell transcriptomes using Iteratively Adjusted-SVA (IA-SVA). <i>Scientific Reports</i> , 2018, 8, 17040. | 1.6 | 8 |
| 16 | V-SVA: an R Shiny application for detecting and annotating hidden sources of variation in single-cell RNA-seq data. <i>Bioinformatics</i> , 2020, 36, 3582-3584. | 1.8 | 7 |