## Noam Auslander

## List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3316835/publications.pdf

Version: 2024-02-01

23 papers

1,549 citations

16 h-index 610901 24 g-index

25 all docs

25 docs citations

25 times ranked

3200 citing authors

| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Robust prediction of response to immune checkpoint blockade therapy in metastatic melanoma. Nature Medicine, 2018, 24, 1545-1549.   | 30.7 | 473       |
| 2  | Genomic determinants of pathogenicity in SARS-CoV-2 and other human coronaviruses. Proceedings of the National Academy of Sciences of the United States of America, 2020, 117, 15193-15199. | 7.1  | 196       |
| 3  | Urea Cycle Dysregulation Generates Clinically Relevant Genomic and Biochemical Signatures. Cell, 2018, 174, 1559-1570.e22.  | 28.9 | 183       |
| 4  | Chemoradiotherapy Resistance in Colorectal Cancer Cells is Mediated by Wnt/ $\hat{l}^2$ -catenin Signaling. Molecular Cancer Research, 2017, 15, 1481-1490.                                 | 3.4  | 105       |
| 5  | Harnessing synthetic lethality to predict the response to cancer treatment. Nature Communications, 2018, 9, 2546.   | 12.8 | 97        |
| 6  | Seeker: alignment-free identification of bacteriophage genomes by deep learning. Nucleic Acids Research, 2020, 48, e121-e121.   | 14.5 | 78        |
| 7  | Co-targeting the tumor endothelium and P-selectin-expressing glioblastoma cells leads to a remarkable therapeutic outcome. ELife, 2017, 6, .  | 6.0  | 50        |
| 8  | Incorporating Machine Learning into Established Bioinformatics Frameworks. International Journal of Molecular Sciences, 2021, 22, 2903.   | 4.1  | 48        |
| 9  | A joint analysis of transcriptomic and metabolomic data uncovers enhanced enzyme-metabolite coupling in breast cancer. Scientific Reports, 2016, 6, 29662.                                  | 3.3  | 43        |
| 10 | An integrated computational and experimental study uncovers <scp>FUT</scp> 9 as a metabolic driver of colorectal cancer. Molecular Systems Biology, 2017, 13, 956.                          | 7.2  | 38        |
| 11 | Pyrvinium Pamoate Induces Death of Triple-Negative Breast Cancer Stem–Like Cells and Reduces Metastases through Effects on Lipid Anabolism. Cancer Research, 2020, 80, 4087-4102.           | 0.9  | 36        |
| 12 | Predicting Complete Remission of Acute Myeloid Leukemia: Machine Learning Applied to Gene Expression. Cancer Informatics, 2019, 18, 117693511983554.  | 1.9  | 27        |
| 13 | A unique insert in the genomes of high-risk human papillomaviruses with a predicted dual role in conferring oncogenic risk. F1000Research, 2019, 8, 1000.                                   | 1.6  | 27        |
| 14 | Interplay between DNA damage repair and apoptosis shapes cancer evolution through aneuploidy and microsatellite instability. Nature Communications, 2020, 11, 1234.                         | 12.8 | 23        |
| 15 | In silico learning of tumor evolution through mutational time series. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 9501-9510.                | 7.1  | 18        |
| 16 | Prediction of the incubation period for COVID-19 and future virus disease outbreaks. BMC Biology, 2020, 18, 186.  | 3.8  | 16        |
| 17 | Reply to: â€`IMPRES does not reproducibly predict response to immune checkpoint blockade therapy in metastatic melanoma'. Nature Medicine, 2019, 25, 1836-1838.                             | 30.7 | 10        |
| 18 | Hard wiring of normal tissue-specific chromosome-wide gene expression levels is an additional factor driving cancer type-specific aneuploidies. Genome Medicine, 2021, 13, 93.              | 8.2  | 10        |

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| 19 | A unique insert in the genomes of high-risk human papillomaviruses with a predicted dual role in conferring oncogenic risk. F1000Research, 2019, 8, 1000.                   | 1.6 | 10       |
| 20 | Single-Cell–Derived Primary Rectal Carcinoma Cell Lines Reflect Intratumor Heterogeneity Associated with Treatment Response. Clinical Cancer Research, 2020, 26, 3468-3480. | 7.0 | 9        |
| 21 | Data-Driven Metabolic Pathway Compositions Enhance Cancer Survival Prediction. PLoS Computational Biology, 2016, 12, e1005125.  | 3.2 | 8        |
| 22 | Identification of combinations of somatic mutations that predict cancer survival and immunotherapy benefit. NAR Cancer, 2021, 3, zcab017.                                   | 3.1 | 2        |
| 23 | The GENDULF algorithm: mining transcriptomics to uncover modifier genes for monogenic diseases.<br>Molecular Systems Biology, 2020, 16, e9701.                              | 7.2 | 2        |