

David Haan

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/9679684/publications.pdf>

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8
papers

686
citations

1684188

5
h-index

1720034

7
g-index

8
all docs

8
docs citations

8
times ranked

1737
citing authors

| # | ARTICLE | IF | CITATIONS |
|---|--|------|-----------|
| 1 | Whole-genome characterization of lung adenocarcinomas lacking alterations in the RTK/RAS/RAF pathway. <i>Cell Reports</i> , 2021, 34, 108707. | 6.4 | 16 |
| 2 | eVIP2: Expression-based variant impact phenotyping to predict the function of gene variants. <i>PLoS Computational Biology</i> , 2021, 17, e1009132. | 3.2 | 8 |
| 3 | A community challenge to evaluate RNA-seq, fusion detection, and isoform quantification methods for cancer discovery. <i>Cell Systems</i> , 2021, 12, 827-838.e5. | 6.2 | 15 |
| 4 | Pathway and network analysis of more than 2500 whole cancer genomes. <i>Nature Communications</i> , 2020, 11, 729. | 12.8 | 73 |
| 5 | Using Transcriptional Signatures to Find Cancer Drivers with LURE. <i>Pacific Symposium on Biocomputing Pacific Symposium on Biocomputing</i> , 2020, 25, 343-354. | 0.7 | 1 |
| 6 | Using Transcriptional Signatures to Find Cancer Drivers with LURE. , 2019, , . | | 3 |
| 7 | Improved genome assembly of American alligator genome reveals conserved architecture of estrogen signaling. <i>Genome Research</i> , 2017, 27, 686-696. | 5.5 | 38 |
| 8 | Comprehensive Molecular Characterization of Pheochromocytoma and Paraganglioma. <i>Cancer Cell</i> , 2017, 31, 181-193. | 16.8 | 532 |