

# Barry Demchak

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/988566/publications.pdf>

Version: 2024-02-01

24  
papers

2,240  
citations

567281  
15  
h-index

642732  
23  
g-index

25  
all docs

25  
docs citations

25  
times ranked

3663  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | Cytoscape Automation: empowering workflow-based network analysis. <i>Genome Biology</i> , 2019, 20, 185.                          | 8.8  | 888       |
| 2  | Using deep learning to model the hierarchical structure and function of a cell. <i>Nature Methods</i> , 2018, 15, 290-298.        | 19.0 | 292       |
| 3  | NDEX, the Network Data Exchange. <i>Cell Systems</i> , 2015, 1, 302-305.  | 6.2  | 247       |
| 4  | Cytoscape: the network visualization tool for GenomeSpace workflows. <i>F1000Research</i> , 2014, 3, 151.                         | 1.6  | 157       |
| 5  | RCy3: Network biology using Cytoscape from within R. <i>F1000Research</i> , 0, 8, 1774.   | 1.6  | 101       |
| 6  | CyREST: Turbocharging Cytoscape Access for External Tools via a RESTful API. <i>F1000Research</i> , 2015, 4, 478.                 | 1.6  | 81        |
| 7  | RCy3: Network biology using Cytoscape from within R. <i>F1000Research</i> , 2019, 8, 1774.  | 1.6  | 80        |
| 8  | Network propagation in the cytoscape cyberinfrastructure. <i>PLoS Computational Biology</i> , 2017, 13, e1005598.                 | 3.2  | 73        |
| 9  | NDEX 2.0: A Clearinghouse for Research on Cancer Pathways. <i>Cancer Research</i> , 2017, 77, e58-e61.                            | 0.9  | 63        |
| 10 | Integrative genomic analysis by interoperation of bioinformatics tools in GenomeSpace. <i>Nature Methods</i> , 2016, 13, 245-247. | 19.0 | 44        |
| 11 | Cytoscape tools for the web age: D3.js and Cytoscape.js exporters. <i>F1000Research</i> , 2014, 3, 143.                           | 1.6  | 44        |
| 12 | RCy3: Network biology using Cytoscape from within R. <i>F1000Research</i> , 2019, 8, 1774.  | 1.6  | 39        |
| 13 | Active Interaction Mapping Reveals the Hierarchical Organization of Autophagy. <i>Molecular Cell</i> , 2017, 65, 761-774.e5.      | 9.7  | 31        |
| 14 | cyNeo4j: connecting Neo4j and Cytoscape. <i>Bioinformatics</i> , 2015, 31, 3868-3869.   | 4.1  | 22        |
| 15 | DDOT: A Swiss Army Knife for Investigating Data-Driven Biological Ontologies. <i>Cell Systems</i> , 2019, 8, 267-273.e3.          | 6.2  | 22        |
| 16 | The Cytoscape Automation app article collection. <i>F1000Research</i> , 2018, 7, 800.   | 1.6  | 17        |
| 17 | The Cytoscape app article collection. <i>F1000Research</i> , 2014, 3, 138.  | 1.6  | 10        |
| 18 | Theory and Application of Network Biology Toward Precision Medicine. <i>Journal of Molecular Biology</i> , 2018, 430, 2873-2874.  | 4.2  | 6         |

| #  | ARTICLE   | IF  | CITATIONS |
|----|---|-----|-----------|
| 19 | aMatReader: Importing adjacency matrices via Cytoscape Automation. <i>F1000Research</i> , 2018, 7, 823. | 1.6 | 6         |
| 20 | aMatReader: Importing adjacency matrices via Cytoscape Automation. <i>F1000Research</i> , 2018, 7, 823. | 1.6 | 5         |
| 21 | Copycat Layout: Network layout alignment via Cytoscape Automation. <i>F1000Research</i> , 2018, 7, 822. | 1.6 | 3         |
| 22 | dot-app: a Graphviz-Cytoscape conversion plug-in. <i>F1000Research</i> , 2016, 5, 2543.                 | 1.6 | 2         |
| 23 | dot-app: a Graphviz-Cytoscape conversion plug-in. <i>F1000Research</i> , 2016, 5, 2543.                 | 1.6 | 2         |
| 24 | Copycat Layout: Network layout alignment via Cytoscape Automation. <i>F1000Research</i> , 2018, 7, 822. | 1.6 | 1         |