

Debra S Goldberg

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/10738134/publications.pdf>

Version: 2024-02-01

20
papers

9,215
citations

687363

13
h-index

839539

18
g-index

22
all docs

22
docs citations

22
times ranked

10381
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Discriminating between HuR and TTP binding sites using the k-spectrum kernel method. PLoS ONE, 2017, 12, e0174052. | 2.5 | 15 |
| 2 | The Topological Profile of a Model of Protein Network Evolution Can Direct Model Improvement. Lecture Notes in Computer Science, 2015, , 40-52. | 1.3 | 2 |
| 3 | Clustering Coefficients in Protein Interaction Hypernetworks. , 2013, , . | | 22 |
| 4 | Evaluating theoretical models of protein interaction network evolution without seed graphs. , 2013, , . | | 0 |
| 5 | Characterization of known protein complexes using k-connectivity and other topological measures. F1000Research, 2013, 2, 172. | 1.6 | 3 |
| 6 | Characterization of known protein complexes using k-connectivity and other topological measures. F1000Research, 2013, 2, 172. | 1.6 | 2 |
| 7 | Improving evolutionary models of protein interaction networks. Bioinformatics, 2011, 27, 376-382. | 4.1 | 24 |
| 8 | Questioning the Ubiquity of Neofunctionalization. PLoS Computational Biology, 2009, 5, e1000252. | 3.2 | 37 |
| 9 | Improving protein function prediction methods with integrated literature data. BMC Bioinformatics, 2008, 9, 198. | 2.6 | 24 |
| 10 | REVERSE ENGINEERING THE EVOLUTION OF PROTEIN INTERACTION NETWORKS. , 2008, , . | | 4 |
| 11 | Predictive models of molecular machines involved in Caenorhabditis elegans early embryogenesis. Nature, 2005, 436, 861-865. | 27.8 | 260 |
| 12 | Towards a proteome-scale map of the human proteinâ€“protein interaction network. Nature, 2005, 437, 1173-1178. | 27.8 | 2,676 |
| 13 | Motifs, themes and thematic maps of an integrated Saccharomyces cerevisiae interaction network. Journal of Biology, 2005, 4, 6. | 2.7 | 154 |
| 14 | Evidence for dynamically organized modularity in the yeast proteinâ€“protein interaction network. Nature, 2004, 430, 88-93. | 27.8 | 1,683 |
| 15 | Global Mapping of the Yeast Genetic Interaction Network. Science, 2004, 303, 808-813. | 12.6 | 1,908 |
| 16 | A Map of the Interactome Network of the Metazoan <i>C. elegans</i>. Science, 2004, 303, 540-543. | 12.6 | 1,587 |
| 17 | Combining biological networks to predict genetic interactions. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 15682-15687. | 7.1 | 225 |
| 18 | Assessing experimentally derived interactions in a small world. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 4372-4376. | 7.1 | 387 |

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 19 | PREY PREFERENCE BY A TOP PREDATOR AND THE STABILITY OF LINKED FOOD CHAINS. <i>Ecology</i> , 2000, 81, 8-14. | 3.2 | 187 |
| 20 | lâ€™m Like You, Just Not In That Way: Tag Networks to Improve Collaborative Filtering. <i>F1000Research</i> , 0, 2, 95. | 1.6 | 0 |