## Bhaskar DasGupta

## List of Publications by Year

 in descending orderSource: https:|/exaly.com/author-pdf/293962/publications.pdf
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Detecting network anomalies using Formanâé"Ricci curvature and a case study for human brain
networks. Scientific Reports, 2021, 11, 8121.

2 A Review of Several Privacy Violation Measures for Large Networks under Active Attacks. , 2020, , .
o
$\begin{array}{lll}3 & \text { A Review of and Some Results for Ollivierâ€"Ricci Network Curvature. Mathematics, 2020, 8, 1416. } & 1.1 \quad 1\end{array}$

4 On theoretical and empirical algorithmic analysis of the efficiency gap measure in partisan
gerrymandering. Journal of Combinatorial Optimization, 2020, 40, 512-546.
$4 \quad \begin{aligned} & \text { On theoretical and empirical algorithmic analysis of the efficiency gap measure } \\ & \text { gerrymandering. Journal of Combinatorial Optimization, 2020, 40, 512-546. }\end{aligned}$
0.8

3

Why Did the Shape of Your Network Change? (On Detecting Network Anomalies via Non-local) Tj ETQq1 $10.784311_{1.0}$ rgBT /Oyerlock

6 On the computational complexities of three problems related to a privacy measure for large networks under active attack. Theoretical Computer Science, 2019, 775, 53-67.
$0.5 \quad 4$

7 A survey of some tensor analysis techniques for biological systems. Quantitative Biology, 2019, 7,
266-277.
0.34
$8 \quad$ On analyzing and evaluating privacy measures for social networks under active attack. Information
Sciences, 2019, 473, 87-100.
4.0

12

9 Effect of Cromov-Hyperbolicity Parameter on Cuts and Expansions in Craphs and Some Algorithmic
Implications. Algorithmica, 2018, 80, 772-800.
1.0

10

10 Topological implications of negative curvature for biological networks. , 2018, , .
0

> On optimal approximability results for computing the strong metric dimension. Discrete Applied
> Mathematics, 2017, 221, 18-24.
0.56

12 Spatio-Temporal Matching for Urban Transportation Applications. ACM Transactions on Spatial Algorithms and Systems, 2017, 3, 1-39.
1.1

7
13 Column-Generation Framework of Nonlinear Similarity Model for Reconstructing Sibling Groups.
$1.0 \quad 5$
INFORMS Journal on Computing, 2015, 27, 35-47.

Algorithmic Perspectives of Network Transitive Reduction Problems and their Applications to
1.3

10
Synthesis and Analysis of Biological Networks. Biology, 2014, 3, 1-21.

Topological implications of negative curvature for biological and social networks. Physical Review E,
0.8

45
2014, 89, 032811.

On the Computational Complexity of Measuring Global Stability of Banking Networks. Algorithmica,
2014, 70, 595-647.
1.0

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17 On a connection between small set expansions and modularity clustering. Information Processing
Letters, 2014, 114, 349-352.
0.4

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19
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A direct variational method for planning monotonically optimal paths for redundant manipulators in
An integrated optimization framework for inferring two generation kinships and parental genotypes
from microsatellite samples. , 2012, , .

On communication protocols that compute almost privately. Theoretical Computer Science, 2012, 457,
$45-58$.

| 25 | Capacitated clustering problem in computational biology: Combinatorial and statistical approach for sibling reconstruction. Computers and Operations Research, 2012, 39, 609-619. | 2.4 | 13 |
| :---: | :---: | :---: | :---: |
| 26 | Models and Algorithmic Tools for Computational Processes in Cellular Biology: Recent Developments and Future Directions. Lecture Notes in Computer Science, 2012, , 84-86. | 1.0 | 0 |
| 27 | Computationally efficient measure of topological redundancy of biological and social networks. Physical Review E, 2011, 84, 036117. | 0.8 | 26 |
| 28 | AN IMPLICIT COVER PROBLEM IN WILD POPULATION STUDY. Discrete Mathematics, Algorithms and Applications, 2010, 02, 21-31. | 0.4 | 2 |

29 COMBINATORIAL RECONSTRUCTION OF HALF-SIBLING GROUPS FROM MICROSATELLITE DATA. Journal of | Bioinformatics and Computational Biology, 2010, 08, 337-356. |
| :--- |
| 12 |

30 Journal on Computing, 2010, 22, 180-194.

1.0

7
31 Inference of Signal Transduction Networks from Double Causal Evidence. Methods in Molecular Biology, 2010, 673, 239-251.
$0.4 \quad 5$

32 A variational approach to path planning for hyper-redundant manipulators. Robotics and Autonomous
3.0

37
Systems, 2009, 57, 194-201.
.
$0.9 \quad 6$
On approximating four covering and packing problems. Journal of Computer and System Sciences,
$232009,75,287-302$.

34 Approximating Transitive Reductions for Directed Networks. Lecture Notes in Computer Science, 2009,
1.0

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Inferring (Biological) Signal Transduction Networks viaÂTransitive Reductions of Directed Graphs.
Algorithmica, 2008, 51, 129-159.
1.0

20
39 Efficient Combinatorial Algorithms for DNA Sequence Processing. , 2007, , 223-239. ..... 040 Reconstructing sibling relationships in wild populations. Bioinformatics, 2007, 23, i49-i56.

| 41 | A Novel Method for Signal Transduction Network Inference from Indirect Experimental Evidence. Journal of Computational Biology, 2007, 14, 927-949. | 0.8 | 52 |
| :---: | :---: | :---: | :---: |
| 42 | Algorithmic and complexity results for decompositions of biological networks into monotone subsystems. BioSystems, 2007, 90, 161-178. | 0.9 | 71 |
| 43 | Randomized approximation algorithms for set multicover problems with applications to reverse engineering of protein and gene networks. Discrete Applied Mathematics, 2007, 155, 733-749. | 0.5 | 35 |
| 44 | Algorithmic Issues in Reverse Engineering of Protein and Gene Networks via the Modular Response Analysis Method. Annals of the New York Academy of Sciences, 2007, 1115, 132-141. | 1.8 | 7 |
| 45 | The inverse protein folding problem on 2D and 3D lattices. Discrete Applied Mathematics, 2007, 155, 719-732. | 0.5 | 7 |

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\begin{aligned}
& \text { Algorithmic and Complexity Results for Decompositions of Biological Networks into Monotone } \\
& \text { Subsystems. Lecture Notes in Computer Science, 2006, , 253-264. }
\end{aligned}
$$

$1.0 \quad 13$

Inapproximability results for the lateral gene transfer problem. Journal of Combinatorial
48 Optimization, 2006, 11, 387-405.
0.8

6

Highly scalable algorithms for robust string barcoding. International Journal of Bioinformatics
$0.1 \quad 16$
Research and Applications, 2005, 1, 145.
$0.1-16$

Tight approximability results for test set problems in bioinformatics. Journal of Computer and System

The Rectangle Enclosure and Point-Dominance Problems Revisited. International Journal of

