

# Philippe Gambette

## List of Publications by Year in descending order

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28  
papers

338  
citations

840776

11  
h-index

839539

18  
g-index

29  
all docs

29  
docs citations

29  
times ranked

332  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Corpus for Idiolectal Research (CIDRE). <i>Journal of Open Humanities Data</i> , 2021, 7, .	0.2	1
2	Valoriser des corpus littéraires numériques avec Wikisource: de la recherche à la pédagogie. , 2021, , 159-176.		1
3	Extracting Event-related Information from a Corpus Regarding Soil Industrial Pollution. , 2021, , .		1
4	Counting phylogenetic networks of level 1 and 2. <i>Journal of Mathematical Biology</i> , 2020, 81, 1357-1395.	1.9	13
5	Normalisation of 16th and 17th century texts in French and geographical named entity recognition. , 2020, , .		5
6	Ce que les données textuelles disent du positionnement en ligne: entre contraintes et appropriations du dispositif. <i>Questions De Communication</i> , 2019, , 123-146.	0.1	3
7	Solving the tree containment problem in linear time for nearly stable phylogenetic networks. <i>Discrete Applied Mathematics</i> , 2018, 246, 62-79.	0.9	13
8	Uprooted Phylogenetic Networks. <i>Bulletin of Mathematical Biology</i> , 2017, 79, 2022-2048.	1.9	6
9	On the challenge of reconstructing level-1 phylogenetic networks from triplets and clusters. <i>Journal of Mathematical Biology</i> , 2017, 74, 1729-1751.	1.9	8
10	Rearrangement moves on rooted phylogenetic networks. <i>PLoS Computational Biology</i> , 2017, 13, e1005611.	3.2	15
11	Development of an in silico method for the identification of subcomplexes involved in the biogenesis of multiprotein complexes in <i>Saccharomyces cerevisiae</i> . <i>BMC Systems Biology</i> , 2017, 11, 67.	3.0	1
12	Do Branch Lengths Help to Locate a Tree in a Phylogenetic Network?. <i>Bulletin of Mathematical Biology</i> , 2016, 78, 1773-1795.	1.9	8
13	Une approche textométrique pour étudier la transmission des savoirs biologiques au XIXe siècle. <i>Nouvelles Perspectives En Sciences Sociales</i> , 2016, 12, 221-253.	0.1	0
14	Locating a Tree in a Phylogenetic Network in Quadratic Time. <i>Lecture Notes in Computer Science</i> , 2015, , 96-107.	1.3	15
15	(Nearly-)tight bounds on the contiguity and linearity of cographs. <i>Theoretical Computer Science</i> , 2014, 522, 1-12.	0.9	4
16	Clust&See: A Cytoscape plugin for the identification, visualization and manipulation of network clusters. <i>BioSystems</i> , 2013, 113, 91-95.	2.0	37
17	Linear-Time Constant-Ratio Approximation Algorithm and Tight Bounds for the Contiguity of Cographs. <i>Lecture Notes in Computer Science</i> , 2013, , 126-136.	1.3	2
18	QUARTETS AND UNROOTED PHYLOGENETIC NETWORKS. <i>Journal of Bioinformatics and Computational Biology</i> , 2012, 10, 1250004.	0.8	29

#	ARTICLE	IF	CITATIONS
19	On encodings of phylogenetic networks of bounded level. <i>Journal of Mathematical Biology</i> , 2012, 65, 157-180.	1.9	27
20	Longueur de branches et arbres de mots. <i>Corpus - Nice</i> , 2012, , .	0.2	1
21	Bootstrap clustering for graph partitioning. <i>RAIRO - Operations Research</i> , 2011, 45, 339-352.	1.8	11
22	Unrestricted and complete Breadth-First Search of trapezoid graphs in time. <i>Information Processing Letters</i> , 2010, 110, 497-502.	0.6	3
23	Visualising a Text with a Tree Cloud. <i>Studies in Classification, Data Analysis, and Knowledge Organization</i> , 2010, , 561-569.	0.2	32
24	Computing galled networks from real data. <i>Bioinformatics</i> , 2009, 25, i85-i93.	4.1	56
25	The Structure of Level- $k$ Phylogenetic Networks. <i>Lecture Notes in Computer Science</i> , 2009, , 289-300.	1.3	14
26	Efficient Neighborhood Encoding for Interval Graphs and Permutation Graphs and $O(n)$ Breadth-First Search. <i>Lecture Notes in Computer Science</i> , 2009, , 146-157.	1.3	4
27	Improved Layout of Phylogenetic Networks. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , 2008, 5, 472-479.	3.0	19
28	On Restrictions of Balanced 2-Interval Graphs. , 2007, , 55-65.		6