

# Nicolas Pasquier

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8178167/publications.pdf>

Version: 2024-02-01

21  
papers

2,431  
citations

1039880

9  
h-index

839398

18  
g-index

22  
all docs

22  
docs citations

22  
times ranked

919  
citing authors

#	ARTICLE	IF	CITATIONS
1	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599.	4.0	9
2	Customer Choice Modelling: A Multi-Level Consensus Clustering Approach. Annals of Emerging Technologies in Computing, 2021, 5, 103-120.	1.0	1
3	A Multi-level Consensus Clustering Framework for Customer Choice Modelling in Travel Industry. Lecture Notes of the Institute for Computer Sciences, Social-Informatics and Telecommunications Engineering, 2020, , 142-157.	0.2	1
4	Using Closed Patterns to Solve the Consensus Clustering Problem. International Journal of Software Engineering and Knowledge Engineering, 2016, 26, 1379-1397.	0.6	3
5	Frequent Closed Patterns Based Multiple Consensus Clustering. Lecture Notes in Computer Science, 2016, , 14-26.	1.0	4
6	Multiple Consensuses Clustering by Iterative Merging/Splitting of Clustering Patterns. Lecture Notes in Computer Science, 2016, , 790-804.	1.0	0
7	The BioKET Biodiversity Data Warehouse: Data and Knowledge Integration and Extraction. Lecture Notes in Computer Science, 2014, , 131-142.	1.0	1
8	A New Approach for Association Rule Mining and Bi-clustering Using Formal Concept Analysis. Lecture Notes in Computer Science, 2012, , 86-101.	1.0	26
9	MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture Notes in Computer Science, 2011, , 174-187.	1.0	5
10	Mining Association Rule Bases from Integrated Genomic Data and Annotations. Lecture Notes in Computer Science, 2009, , 78-90.	1.0	1
11	GenMiner: mining non-redundant association rules from integrated gene expression data and annotations. Bioinformatics, 2008, 24, 2643-2644.	1.8	56
12	GenMiner: Mining Informative Association Rules from Genomic Data. , 2007, , .		13
13	Co-expressed gene groups analysis (CGGA): An automatic tool for the interpretation of microarray experiments. Journal of Integrative Bioinformatics, 2006, 3, 188-198.	1.0	2
14	Interpreting Microarray Experiments Via Co-expressed Gene Groups Analysis (CGGA). Lecture Notes in Computer Science, 2006, , 316-320.	1.0	0
15	Generating a Condensed Representation for Association Rules. Journal of Intelligent Information Systems, 2005, 24, 29-60.	2.8	154
16	Computing iceberg concept lattices with Titanic. Data and Knowledge Engineering, 2002, 42, 189-222.	2.1	336
17	Intelligent Structuring and Reducing of Association Rules with Formal Concept Analysis. Lecture Notes in Computer Science, 2001, , 335-350.	1.0	58
18	Mining frequent patterns with counting inference. SIGKDD Explorations: Newsletter of the Special Interest Group (SIG) on Knowledge Discovery & Data Mining, 2000, 2, 66-75.	3.2	217

#	ARTICLE	IF	CITATIONS
19	Mining Minimal Non-redundant Association Rules Using Frequent Closed Itemsets. Lecture Notes in Computer Science, 2000, , 972-986.	1.0	187
20	Discovering Frequent Closed Itemsets for Association Rules. Lecture Notes in Computer Science, 1999, , 398-416.	1.0	740
21	Efficient mining of association rules using closed itemset lattices. Information Systems, 1999, 24, 25-46.	2.4	609