Nicolas Pasquier

List of Publications by Year in descending order

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1039880 839398 2,431 21 9 18 citations h-index g-index papers 22 22 22 919 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Discovering Frequent Closed Itemsets for Association Rules. Lecture Notes in Computer Science, 1999, , 398-416.	1.0	740
2	Efficient mining of association rules using closed itemset lattices. Information Systems, 1999, 24, 25-46.	2.4	609
3	Computing iceberg concept lattices with Titanic. Data and Knowledge Engineering, 2002, 42, 189-222.	2.1	336
4	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
5	Mining Minimal Non-redundant Association Rules Using Frequent Closed Itemsets. Lecture Notes in Computer Science, 2000, , 972-986.	1.0	187
6	Generating a Condensed Representation for Association Rules. Journal of Intelligent Information Systems, 2005, 24, 29-60.	2.8	154
7	Intelligent Structuring and Reducing of Association Rules with Formal Concept Analysis. Lecture Notes in Computer Science, 2001, , 335-350.	1.0	58
8	GenMiner: mining non-redundant association rules from integrated gene expression data and annotations. Bioinformatics, 2008, 24, 2643-2644.	1.8	56
9	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
10	GenMiner: Mining Informative Association Rules from Genomic Data., 2007,,.		13
10	GenMiner: Mining Informative Association Rules from Genomic Data., 2007,,. Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599.	4.0	13
	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235,	4.0	
11	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599. MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture		9
11 12	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599. MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture Notes in Computer Science, 2011, , 174-187. Frequent Closed Patterns Based Multiple Consensus Clustering. Lecture Notes in Computer Science,	1.0	5
11 12 13	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599. MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture Notes in Computer Science, 2011, , 174-187. Frequent Closed Patterns Based Multiple Consensus Clustering. Lecture Notes in Computer Science, 2016, , 14-26. Using Closed Patterns to Solve the Consensus Clustering Problem. International Journal of Software	1.0	9 5 4
11 12 13	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599. MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture Notes in Computer Science, 2011, , 174-187. Frequent Closed Patterns Based Multiple Consensus Clustering. Lecture Notes in Computer Science, 2016, , 14-26. Using Closed Patterns to Solve the Consensus Clustering Problem. International Journal of Software Engineering and Knowledge Engineering, 2016, 26, 1379-1397. Co-expressed gene groups analysis (CGGA): An automatic tool for the interpretation of microarray	1.0	9 5 4 3
11 12 13 14	Semi-supervised consensus clustering based on closed patterns. Knowledge-Based Systems, 2022, 235, 107599. MOSCFRA: A Multi-objective Genetic Approach for Simultaneous Clustering and Gene Ranking. Lecture Notes in Computer Science, 2011, , 174-187. Frequent Closed Patterns Based Multiple Consensus Clustering. Lecture Notes in Computer Science, 2016, , 14-26. Using Closed Patterns to Solve the Consensus Clustering Problem. International Journal of Software Engineering and Knowledge Engineering, 2016, 26, 1379-1397. Co-expressed gene groups analysis (CGGA): An automatic tool for the interpretation of microarray experiments. Journal of Integrative Bioinformatics, 2006, 3, 188-198. Customer Choice Modelling: A Multi-Level Consensus Clustering Approach. Annals of Emerging	1.0 1.0 0.6	9 5 4 3

#	Article	IF	CITATIONS
19	The BioKET Biodiversity Data Warehouse: Data and Knowledge Integration and Extraction. Lecture Notes in Computer Science, 2014, , 131-142.	1.0	1
20	Interpreting Microarray Experiments Via Co-expressed Gene Groups Analysis (CGGA). Lecture Notes in Computer Science, 2006, , 316-320.	1.0	0
21	Multiple Consensuses Clustering by Iterative Merging/Splitting of Clustering Patterns. Lecture Notes in Computer Science, 2016, , 790-804.	1.0	O