## Erhard Rahm

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3671767/publications.pdf

Version: 2024-02-01

172207 76769 7,988 161 29 74 citations h-index g-index papers 177 177 177 4065 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	A survey of approaches to automatic schema matching. VLDB Journal, 2001, 10, 334-350.	2.7	2,288
2	COMA — A system for flexible combination of schema matching approaches. , 2002, , 610-621.		580
3	Schema and ontology matching with COMA++., 2005,,.		395
4	Frameworks for entity matching: A comparison. Data and Knowledge Engineering, 2010, 69, 197-210.	2.1	294
5	Evaluation of entity resolution approaches on real-world match problems. Proceedings of the VLDB Endowment, 2010, 3, 484-493.	2.1	275
6	AgentWork: a workflow system supporting rule-based workflow adaptation. Data and Knowledge Engineering, 2004, 51, 223-256.	2.1	241
7	Matching large schemas: Approaches and evaluation. Information Systems, 2007, 32, 857-885.	2.4	183
8	FUNC: a package for detecting significant associations between gene sets and ontological annotations. BMC Bioinformatics, 2007, 8, 41.	1.2	180
9	Rondo., 2003,,.		175
10	Comparison of Schema Matching Evaluations. Lecture Notes in Computer Science, 2003, , 221-237.	1.0	150
11	Generic schema matching, ten years later. Proceedings of the VLDB Endowment, 2011, 4, 695-701.	2.1	139
12	A survey of current Link Discovery frameworks. Semantic Web, 2016, 8, 419-436.	1.1	131
13	Dedoop. Proceedings of the VLDB Endowment, 2012, 5, 1878-1881.	2.1	107
14	Load Balancing for MapReduce-based Entity Resolution. , 2012, , .		101
15	Towards Large-Scale Schema and Ontology Matching. , 2011, , 3-27.		100
16	Smart Medical Information Technology for Healthcare (SMITH). Methods of Information in Medicine, 2018, 57, e92-e105.	0.7	89
17	Matching large XML schemas. SIGMOD Record, 2004, 33, 26-31.	0.7	76
18	Convergent validity of bibliometric Google Scholar data in the field of chemistryâ€"Citation counts for papers that were accepted by Angewandte Chemie International Edition or rejected but published elsewhere, using Google Scholar, Science Citation Index, Scopus, and Chemical Abstracts. Journal of Informetrics, 2009, 3, 27-35.	1.4	76

#	Article	IF	CITATIONS
19	Privacy-Preserving Record Linkage for Big Data: Current Approaches and Research Challenges. , 2017, , 851-895.		73
20	An online bibliography on schema evolution. SIGMOD Record, 2006, 35, 30-31.	0.7	65
21	Citation analysis of database publications. SIGMOD Record, 2005, 34, 48-53.	0.7	64
22	MediGRID: Towards a user friendly secured grid infrastructure. Future Generation Computer Systems, 2009, 25, 326-336.	4.9	63
23	COnto–Diff: generation of complex evolution mappings for life science ontologies. Journal of Biomedical Informatics, 2013, 46, 15-32.	2.5	60
24	Multi-pass sorted neighborhood blocking with MapReduce. Computer Science - Research and Development, 2012, 27, 45-63.	2.7	58
25	Enriching ontology mappings with semantic relations. Data and Knowledge Engineering, 2014, 93, 1-18.	2.1	58
26	Supporting executable mappings in model management. , 2005, , .		56
27	Recent Advances in Schema and Ontology Evolution. , 2011, , 149-190.		54
28	XMach-1: A Benchmark for XML Data Management. Informatik Aktuell, 2001, , 264-273.	0.4	53
29	Empirical performance evaluation of concurrency and coherency control protocols for database sharing systems. ACM Transactions on Database Systems, 1993, 18, 333-377.	1.5	51
30	Quickmig. , 2007, , .		50
31	GOMMA: a component-based infrastructure for managing and analyzing life science ontologies and their evolution. Journal of Biomedical Semantics, 2011, 2, 6.	0.9	49
32	Tailoring entity resolution for matching product offers. , 2012, , .		49
33	Data Warehouse Scenarios for Model Management. Lecture Notes in Computer Science, 2000, , 1-15.	1.0	41
34	AMC - A framework for modelling and comparing matching systems as matching processes. , 2011, , .		37
35	Using Link Features for Entity Clustering in Knowledge Graphs. Lecture Notes in Computer Science, 2018, , 576-592.	1.0	37
36	A Clustering-Based Framework to Control Block Sizes for Entity Resolution. , 2015, , .		37

#	Article	IF	Citations
37	Management and Analysis of Big Graph Data: Current Systems and Open Challenges., 2017,, 457-505.		36
38	On Matching Large Life Science Ontologies in Parallel. Lecture Notes in Computer Science, 2010, , 35-49.	1.0	36
39	ATOM: Automatic target-driven ontology merging. , 2011, , .		35
40	Impact of ontology evolution on functional analyses. Bioinformatics, 2012, 28, 2671-2677.	1.8	33
41	Evolution of biomedical ontologies and mappings: Overview of recent approaches. Computational and Structural Biotechnology Journal, 2016, 14, 333-340.	1.9	33
42	The Case for Holistic Data Integration. Lecture Notes in Computer Science, 2016, , 11-27.	1.0	32
43	Parallel Entity Resolution with Dedoop. Datenbank-Spektrum, 2013, 13, 23-32.	1.2	31
44	Target-driven merging of taxonomies with Atom. Information Systems, 2014, 42, 1-14.	2.4	29
45	A Framework for workload allocation in distributed transaction processing systems. Journal of Systems and Software, 1992, 18, 171-190.	3.3	28
46	Rewrite techniques for performance optimization of schema matching processes. , 2010, , .		28
47	Comparative Evaluation of Distributed Clustering Schemes for Multi-source Entity Resolution. Lecture Notes in Computer Science, 2017, , 278-293.	1.0	27
48	Towards a Benchmark for Ontology Merging. Lecture Notes in Computer Science, 2012, , 124-133.	1.0	27
49	Learning-Based Approaches for Matching Web Data Entities. IEEE Internet Computing, 2010, 14, 23-31.	3.2	26
50	Comparative evaluation of entity resolution approaches with FEVER. Proceedings of the VLDB Endowment, 2009, 2, 1574-1577.	2.1	25
51	A Self-Configuring Schema Matching System. , 2012, , .		25
52	Primary copy synchronization for DB-Sharing. Information Systems, 1986, 11, 275-286.	2.4	24
53	OnEX: Exploring changes in life science ontologies. BMC Bioinformatics, 2009, 10, 250.	1.2	23
54	Analyzing the Evolution of Life Science Ontologies and Mappings. Lecture Notes in Computer Science, 2008, , 11-27.	1.0	23

#	Article	IF	CITATIONS
55	The Scholarly Impact of CLEF (2000–2009). Lecture Notes in Computer Science, 2013, , 1-12.	1.0	23
56	Parallel query processing in shared disk database systems. SIGMOD Record, 1993, 22, 32-37.	0.7	22
57	Developing metadata-intensive applications with Rondo. Web Semantics, 2003, 1, 47-74.	2.2	21
58	Evaluation of metadata representations in RDF stores. Semantic Web, 2019, 10, 205-229.	1.1	21
59	Dynamic load balancing in parallel database systems. Lecture Notes in Computer Science, 1996, , 37-52.	1.0	20
60	Analyzing extended property graphs with Apache Flink. , 2016, , .		20
61	Instance-Based Matching of Large Life Science Ontologies. , 2007, , 172-187.		20
62	Declarative and distributed graph analytics with GRADOOP. Proceedings of the VLDB Endowment, 2018, 11, 2006-2009.	2.1	19
63	Block-based load balancing for entity resolution with MapReduce. , 2011, , .		18
64	Holistic Entity Clustering for Linked Data., 2016,,.		18
64 65	Holistic Entity Clustering for Linked Data., 2016,,.  Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004,, 811-822.	1.0	18
	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture	0.5	
65	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004, , 811-822.  Scalable Matching and Clustering of Entities with FAMER. Complex Systems Informatics and Modeling		18
66	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004, , 811-822.  Scalable Matching and Clustering of Entities with FAMER. Complex Systems Informatics and Modeling Quarterly, 2018, , 61-83.  A multi-part matching strategy for mapping LOINC with laboratory terminologies. Journal of the	0.5	18
65 66 67	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004, , 811-822.  Scalable Matching and Clustering of Entities with FAMER. Complex Systems Informatics and Modeling Quarterly, 2018, , 61-83.  A multi-part matching strategy for mapping LOINC with laboratory terminologies. Journal of the American Medical Informatics Association: JAMIA, 2014, 21, 792-800.  Graph-based data integration and business intelligence with BIIIG. Proceedings of the VLDB	0.5	18 18 17
65 66 67 68	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004, , 811-822.  Scalable Matching and Clustering of Entities with FAMER. Complex Systems Informatics and Modeling Quarterly, 2018, , 61-83.  A multi-part matching strategy for mapping LOINC with laboratory terminologies. Journal of the American Medical Informatics Association: JAMIA, 2014, 21, 792-800.  Graph-based data integration and business intelligence with BIIIG. Proceedings of the VLDB Endowment, 2014, 7, 1577-1580.	0.5	18 18 17 17
65 66 67 68	Flexible Integration of Molecular-Biological Annotation Data: The GenMapper Approach. Lecture Notes in Computer Science, 2004, , 811-822.  Scalable Matching and Clustering of Entities with FAMER. Complex Systems Informatics and Modeling Quarterly, 2018, , 61-83.  A multi-part matching strategy for mapping LOINC with laboratory terminologies. Journal of the American Medical Informatics Association: JAMIA, 2014, 21, 792-800.  Graph-based data integration and business intelligence with BIIIG. Proceedings of the VLDB Endowment, 2014, 7, 1577-1580.  Extracting Semantic Concept Relations from Wikipedia. , 2014, , .	0.5 2.2 2.1	18 18 17 17 16

#	Article	IF	Citations
73	Affiliation analysis of database publications. SIGMOD Record, 2011, 40, 26-31.	0.7	14
74	CODEX: exploration of semantic changes between ontology versions. Bioinformatics, 2012, 28, 895-896.	1.8	14
75	Don't match twice. , 2013, , .		14
76	Scalable Privacy-Preserving Linking of Multiple Databases Using Counting Bloom Filters. , 2016, , .		14
77	Cypher-based Graph Pattern Matching in Gradoop. , 2017, , .		14
78	Semi-automatic Adaptation of Mappings between Life Science Ontologies. Lecture Notes in Computer Science, 2013, , 90-104.	1.0	14
79	Parallel Privacy-preserving Record Linkage using LSH-based Blocking. , 2018, , .		14
80	Incremental Clustering on Linked Data., 2018,,.		13
81	Incremental clustering techniques for multi-party Privacy-Preserving Record Linkage. Data and Knowledge Engineering, 2020, 128, 101809.	2.1	13
82	Distributed temporal graph analytics with GRADOOP. VLDB Journal, 2022, 31, 375-401.	2.7	13
83	Leveraging the Impact of Ontology Evolution on Semantic Annotations. Lecture Notes in Computer Science, 2016, , 68-82.	1.0	12
84	Learning-based entity resolution with MapReduce. , 2011, , .		11
85	Iterative Computation of Connected Graph Components with MapReduce. Datenbank-Spektrum, 2014, 14, 107-117.	1.2	11
86	BioFuice: Mapping-Based Data Integration in Bioinformatics. Lecture Notes in Computer Science, 2006, , 124-135.	1.0	11
87	Incremental Multi-source Entity Resolution for Knowledge Graph Completion. Lecture Notes in Computer Science, 2020, , 393-408.	1.0	11
88	When to Reach for the Cloud: Using Parallel Hardware for Link Discovery. Lecture Notes in Computer Science, 2013, , 275-289.	1.0	11
89	BIIIG: Enabling business intelligence with integrated instance graphs. , 2014, , .		10
90	Performance evaluation of parallel transaction processing in shared nothing database systems. Lecture Notes in Computer Science, 1992, , 295-310.	1.0	10

#	Article	IF	CITATIONS
91	Efficient Management of Biomedical Ontology Versions. Lecture Notes in Computer Science, 2009, , 574-583.	1.0	9
92	Automatic Extraction of Semantic Relations from Wikipedia. International Journal on Artificial Intelligence Tools, 2015, 24, 1540010.	0.7	9
93	Speeding up Privacy Preserving Record Linkage for Metric Space Similarity Measures. Datenbank-Spektrum, 2016, 16, 227-236.	1.2	9
94	DIMSpan., 2017,,.		9
95	Estimating the Quality of Ontology-Based Annotations by Considering Evolutionary Changes. Lecture Notes in Computer Science, 2009, , 71-87.	1.0	9
96	Semantic Enrichment of Ontology Mappings: A Linguistic-Based Approach. Lecture Notes in Computer Science, 2013, , 42-55.	1.0	9
97	WETSUIT. Proceedings of the VLDB Endowment, 2012, 5, 1970-1973.	2.1	9
98	A Reuse-Based Annotation Approach for Medical Documents. Lecture Notes in Computer Science, 2016, , 135-150.	1.0	8
99	FoodBroker - Generating Synthetic Datasets for Graph-Based Business Analytics. Lecture Notes in Computer Science, 2015, , 145-155.	1.0	8
100	Dynamic query scheduling in parallel data warehouses. Concurrency Computation Practice and Experience, 2003, 15, 1169-1190.	1.4	7
101	Entity Search Strategies for Mashup Applications. , 2012, , .		7
102	Distributed Holistic Clustering on Linked Data. Lecture Notes in Computer Science, 2017, , 371-382.	1.0	7
103	Graph Mining for Complex Data Analytics. , 2016, , .		6
104	Mining and ranking of generalized multi-dimensional frequent subgraphs. , 2017, , .		6
105	ScaDS Research on Scalable Privacy-preserving Record Linkage. Datenbank-Spektrum, 2019, 19, 31-40.	1.2	6
106	Post-processing Methods for High Quality Privacy-Preserving Record Linkage. Lecture Notes in Computer Science, 2018, , 263-278.	1.0	6
107	Annotating Medical Forms Using UMLS. Lecture Notes in Computer Science, 2015, , 55-69.	1.0	6
108	Dynamic Fusion of Web Data. Lecture Notes in Computer Science, 2007, , 14-16.	1.0	6

#	Article	IF	CITATIONS
109	Analysis of parallel scan processing in Shared Disk database systems. Lecture Notes in Computer Science, 1995, , 485-500.	1.0	5
110	Big Data Analytics. IT - Information Technology, 2016, 58, 155-156.	0.6	5
111	Analyzing Temporal Graphs with Gradoop. Datenbank-Spektrum, 2019, 19, 199-208.	1.2	5
112	Big Data Competence Center ScaDS Dresden/Leipzig: Overview and selected research activities. Datenbank-Spektrum, 2019, 19, 5-16.	1.2	5
113	Discovering Evolving Regions in Life Science Ontologies. Lecture Notes in Computer Science, 2010, , 19-34.	1.0	5
114	Effective Composition of Mappings for Matching Biomedical Ontologies. Lecture Notes in Computer Science, 2015, , 176-190.	1.0	5
115	PRIMAT. Proceedings of the VLDB Endowment, 2019, 12, 1826-1829.	2.1	5
116	Automatic Optimization of Web Recommendations Using Feedback and Ontology Graphs. Lecture Notes in Computer Science, 2005, , 375-386.	1.0	4
117	Management of evolving semantic grid metadata within a collaborative platform. Information Sciences, 2010, 180, 1837-1849.	4.0	4
118	PDFMeat., 2011,,.		4
119	Discovering product counterfeits in online shops. Journal of Data and Information Quality, 2014, 5, 1-3.	1.5	4
120	Distributed Privacy-Preserving Record Linkage Using Pivot-Based Filter Techniques. , 2018, , .		4
121	Optimization of the Mainzelliste software for fast privacy-preserving record linkage. Journal of Translational Medicine, 2021, 19, 33.	1.8	4
122	An Integrated Platform for Analyzing Molecular-Biological Data Within Clinical Studies. Lecture Notes in Computer Science, 2006, , 399-410.	1.0	4
123	Evaluating and Improving Annotation Tools for Medical Forms. Lecture Notes in Computer Science, 2017, , 1-16.	1.0	4
124	Evaluating Cross-lingual Semantic Annotation for Medical Forms. , 2020, , .		4
125	LEAPME: Learning-based Property Matching with Embeddings. Data and Knowledge Engineering, 2022, 137, 101943.	2.1	4
126	Adaptive website recommendations with AWESOME. VLDB Journal, 2005, 14, 357-372.	2.7	3

#	Article	IF	CITATIONS
127	Rule-based construction of matching processes. , 2011, , .		3
128	Hybrid Integration of Molecular-Biological Annotation Data. Lecture Notes in Computer Science, 2005, , 208-223.	1.0	3
129	Datenbanksysteme., 1999,,.		3
130	Holistic Schema Matching. , 2019, , 960-965.		3
131	Performance evaluation of extended storage architectures for transaction processing. SIGMOD Record, 1992, 21, 308-317.	0.7	2
132	Semi-Automatic Identification of Counterfeit Offers in Online Shopping Platforms. Journal of Internet Commerce, 2016, 15, 59-75.	3.5	2
133	Evolving semantic annotations through multiple versions of controlled medical terminologies. Health and Technology, 2018, 8, 361-376.	2.1	2
134	BIGGR: Bringing Gradoop to Applications. Datenbank-Spektrum, 2019, 19, 51-60.	1.2	2
135	Evolution Analysis of Large Graphs with Gradoop. Communications in Computer and Information Science, 2020, , 402-408.	0.4	2
136	Large-Scale Schema Matching. , 2018, , 1-6.		2
137	A Platform for Collaborative Management of Semantic Grid Metadata. Studies in Computational Intelligence, 2008, , 115-125.	0.7	2
138	Evaluierung von Data Warehouse-Werkzeugen. , 2000, , 43-57.		2
139	Concurrency Control in DB-Sharing Systems. Informatik-Fachberichte, 1986, , 617-632.	0.2	2
140	Combining Semantic and Lexical Measures to Evaluate Medical Terms Similarity. Lecture Notes in Computer Science, 2019, , 17-32.	1.0	2
141	Large-Scale Entity Resolution. , 2019, , 1100-1105.		2
142	High performance cache management for sequential data access. , 1992, , .		1
143	Enhancing Cross-lingual Semantic Annotations using Deep Network Sentence Embeddings., 2021,,.		1
144	Restricting the overlap of Top-N sets in schema matching. , 2011, , .		1

#	Article	IF	CITATIONS
145	Developing Metadata-Intensive Applications with Rondo. SSRN Electronic Journal, 0, , .	0.4	1
146	Holistic Schema Matching., 2018, , 1-5.		1
147	A Learning-Based Approach to Combine Medical Annotation Results. Lecture Notes in Computer Science, 2019, , 135-143.	1.0	1
148	Informativeness-Based Active Learning for Entity Resolution. Communications in Computer and Information Science, 2020, , 125-141.	0.4	1
149	ERGAN: Generative Adversarial Networks for Entity Resolution. , 2020, , .		1
150	High performance cache management for sequential data access. Performance Evaluation Review, 1992, 20, 243-244.	0.4	0
151	Cache management for shared sequential data access. Information Systems, 1993, 18, 197-213.	2.4	O
152	Architekturen für verteiltes und paralleles Datenmanagement. EXamen Press, 2015, , 43-78.	0.0	0
153	ScaDS Dresden/Leipzig – A competence center for collaborative big data research. IT - Information Technology, 2018, 60, 327-333.	0.6	O
154	Towards the smart use of embedding and instance features for property matching. , 2021, , .		0
155	A Reliable and Efficient Synchronization Protocol for Database Sharing Systems. Informatik-Fachberichte, 1987, , 336-347.	0.2	0
156	Mehrrechner-Datenbanksysteme. Informatik-Fachberichte, 1988, , 6-19.	0.2	0
157	ArchitekturansÃtze zur Unterstützung heterogener Datenbanken. Informatik Aktuell, 1992, , 106-118.	0.4	O
158	Towards a Multi-level Approach for the Maintenance of Semantic Annotations. , 2017, , .		0
159	Large-Scale Schema Matching. , 2019, , 1105-1110.		0
160	Parameterized XPath Views. Lecture Notes in Computer Science, 2007, , 125-137.	1.0	0
161	Multi-source dataset of e-commerce products with attributes for property matching. Data in Brief, 2022, 41, 107884.	0.5	0