

# Xiaoqian Jiang

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

90  
papers

2,523  
citations

26  
h-index

48  
g-index

117  
ext. papers

3,487  
ext. citations

6.4  
avg, IF

5.53  
L-index

#	Paper	IF	Citations
90	Factors Associated With COVID-19 Death in the United States: Cohort Study.. <i>JMIR Public Health and Surveillance</i> , <b>2022</b> ,	11.4	2
89	Counterfactual analysis of differential comorbidity risk factors in Alzheimer's disease and related dementias <b>2022</b> , 1, e0000018		0
88	Privacy-preserving logistic regression with secret sharing.. <i>BMC Medical Informatics and Decision Making</i> , <b>2022</b> , 22, 89	3.6	1
87	Deep graph convolutional network for US birth data harmonization.. <i>Journal of Biomedical Informatics</i> , <b>2021</b> , 125, 103974	10.2	
86	Drug repurposing for COVID-19 using graph neural network and harmonizing multiple evidence. <i>Scientific Reports</i> , <b>2021</b> , 11, 23179	4.9	5
85	Deep representation learning of patient data from Electronic Health Records (EHR): A systematic review. <i>Journal of Biomedical Informatics</i> , <b>2021</b> , 115, 103671	10.2	10
84	COVID-19 trial graph: a linked graph for COVID-19 clinical trials. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2021</b> , 28, 1964-1969	8.6	1
83	Human Endogenous Retroviruses in Glioblastoma Multiforme. <i>Microorganisms</i> , <b>2021</b> , 9,	4.9	6
82	Privacy-protecting, reliable response data discovery using COVID-19 patient observations. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2021</b> , 28, 1765-1776	8.6	3
81	Calibrating predictive model estimates in a distributed network of patient data. <i>Journal of Biomedical Informatics</i> , <b>2021</b> , 117, 103758	10.2	0
80	Hyperpolarized Magnetic Resonance and Artificial Intelligence: Frontiers of Imaging in Pancreatic Cancer. <i>JMIR Medical Informatics</i> , <b>2021</b> , 9, e26601	3.6	1
79	Are synthetic clinical notes useful for real natural language processing tasks: A case study on clinical entity recognition. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2021</b> , 28, 2193-2201	8.6	3
78	Contact Tracing Apps: Lessons Learned on Privacy, Autonomy, and the Need for Detailed and Thoughtful Implementation. <i>JMIR Medical Informatics</i> , <b>2021</b> , 9, e27449	3.6	3
77	Anticancer drug synergy prediction in understudied tissues using transfer learning. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2021</b> , 28, 42-51	8.6	16
76	Privacy-preserving string search on encrypted genomic data using a generalized suffix tree. <i>Informatics in Medicine Unlocked</i> , <b>2021</b> , 23, 100525	5.3	1
75	Demystifying the Dark Web Opioid Trade: Content Analysis on Anonymous Market Listings and Forum Posts. <i>Journal of Medical Internet Research</i> , <b>2021</b> , 23, e24486	7.6	1
74	Ultrafast homomorphic encryption models enable secure outsourcing of genotype imputation. <i>Cell Systems</i> , <b>2021</b> , 12, 1108-1120.e4	10.6	4

73	Noise-tolerant similarity search in temporal medical data. <i>Journal of Biomedical Informatics</i> , <b>2021</b> , 113, 103667	10.2	1
72	Harmonized representation learning on dynamic EHR graphs. <i>Journal of Biomedical Informatics</i> , <b>2020</b> , 106, 103426	10.2	7
71	COVID-19 TestNorm: A tool to normalize COVID-19 testing names to LOINC codes. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2020</b> , 27, 1437-1442	8.6	8
70	Temporal phenotyping for transitional disease progress: An application to epilepsy and Alzheimer's disease. <i>Journal of Biomedical Informatics</i> , <b>2020</b> , 107, 103462	10.2	1
69	VERTICOX: Vertically Distributed Cox Proportional Hazards Model Using the Alternating Direction Method of Multipliers. <i>IEEE Transactions on Knowledge and Data Engineering</i> , <b>2020</b> , 1-1	4.2	4
68	Predict or draw blood: An integrated method to reduce lab tests. <i>Journal of Biomedical Informatics</i> , <b>2020</b> , 104, 103394	10.2	4
67	Multimodal Phenotyping of Alzheimer's Disease with Longitudinal Magnetic Resonance Imaging and Cognitive Function Data. <i>Scientific Reports</i> , <b>2020</b> , 10, 5527	4.9	7
66	Drug Repurposing for COVID-19 using Graph Neural Network with Genetic, Mechanistic, and Epidemiological Validation <b>2020</b> ,		13
65	A deep learning solution to recommend laboratory reduction strategies in ICU. <i>International Journal of Medical Informatics</i> , <b>2020</b> , 144, 104282	5.3	1
64	SCOR: A secure international informatics infrastructure to investigate COVID-19. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2020</b> , 27, 1721-1726	8.6	20
63	Multiple imputation for analysis of incomplete data in distributed health data networks. <i>Nature Communications</i> , <b>2020</b> , 11, 5467	17.4	6
62	A secure system for genomics clinical decision support. <i>Journal of Biomedical Informatics</i> , <b>2020</b> , 112, 103602	10.2	2
61	Model-Protected Multi-Task Learning. <i>IEEE Transactions on Pattern Analysis and Machine Intelligence</i> , <b>2020</b> , PP,	13.3	1
60	Identification of de novo mutations in prenatal neurodevelopment-associated genes in schizophrenia in two Han Chinese patient-sibling family-based cohorts. <i>Translational Psychiatry</i> , <b>2020</b> , 10, 307	8.6	1
59	Treating medical data as a durable asset. <i>Nature Genetics</i> , <b>2020</b> , 52, 1005-1010	36.3	5
58	Big Data Privacy in Biomedical Research. <i>IEEE Transactions on Big Data</i> , <b>2020</b> , 6, 296-308	3.2	11
57	Distributed learning from multiple EHR databases: Contextual embedding models for medical events. <i>Journal of Biomedical Informatics</i> , <b>2019</b> , 92, 103138	10.2	15
56	A Predictive Model for Determining Patients Not Requiring Prolonged Hospital Length of Stay After Elective Primary Total Hip Arthroplasty. <i>Anesthesia and Analgesia</i> , <b>2019</b> , 129, 43-50	3.9	16

55	Privacy-preserving techniques of genomic data-a survey. <i>Briefings in Bioinformatics</i> , <b>2019</b> , 20, 887-895	13.4	19
54	SAFETY: Secure gwAs in Federated Environment through a hYbrid Solution. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , <b>2019</b> , 16, 93-102	3	21
53	Patient ranking with temporally annotated data. <i>Journal of Biomedical Informatics</i> , <b>2018</b> , 78, 43-53	10.2	6
52	Selecting Optimal Subset to release under Differentially Private M-estimators from Hybrid Datasets. <i>IEEE Transactions on Knowledge and Data Engineering</i> , <b>2018</b> , 30, 573-584	4.2	5
51	Deep learning for healthcare: review, opportunities and challenges. <i>Briefings in Bioinformatics</i> , <b>2018</b> , 19, 1236-1246	13.4	762
50	Privacy-Preserving Patient Similarity Learning in a Federated Environment: Development and Analysis. <i>JMIR Medical Informatics</i> , <b>2018</b> , 6, e20	3.6	53
49	Secure Logistic Regression Based on Homomorphic Encryption: Design and Evaluation. <i>JMIR Medical Informatics</i> , <b>2018</b> , 6, e19	3.6	71
48	Privacy-Preserving Predictive Modeling: Harmonization of Contextual Embeddings From Different Sources. <i>JMIR Medical Informatics</i> , <b>2018</b> , 6, e33	3.6	8
47	Secure and Efficient Regression Analysis Using a Hybrid Cryptographic Framework: Development and Evaluation. <i>JMIR Medical Informatics</i> , <b>2018</b> , 6, e14	3.6	4
46	GenoPri'16: International Workshop on Genome Privacy and Security. <i>IEEE/ACM Transactions on Computational Biology and Bioinformatics</i> , <b>2018</b> , 15, 1403-1404	3	
45	iDASH secure genome analysis competition 2017. <i>BMC Medical Genomics</i> , <b>2018</b> , 11, 85	3.7	9
44	Secure Outsourced Matrix Computation and Application to Neural Networks <b>2018</b> , 2018, 1209-1222		62
43	Privacy Policy and Technology in Biomedical Data Science. <i>Annual Review of Biomedical Data Science</i> , <b>2018</b> , 1, 115-129	5.6	11
42	PRINCESS: Privacy-protecting Rare disease International Network Collaboration via Encryption through Software guard extensionS. <i>Bioinformatics</i> , <b>2017</b> , 33, 871-878	7.2	36
41	Discriminative and Distinct Phenotyping by Constrained Tensor Factorization. <i>Scientific Reports</i> , <b>2017</b> , 7, 1114	4.9	14
40	Partitioning-based mechanisms under personalized differential privacy. <i>Lecture Notes in Computer Science</i> , <b>2017</b> , 10234, 615-627	0.9	10
39	Addressing Beacon re-identification attacks: quantification and mitigation of privacy risks. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2017</b> , 24, 799-805	8.6	43
38	A community effort to protect genomic data sharing, collaboration and outsourcing. <i>Npj Genomic Medicine</i> , <b>2017</b> , 2, 33	6.2	22

37	DiagTree <b>2017</b> ,		2
36	PRESAGE: PRivacy-preserving gEnetic testing via SoftwAre Guard Extension. <i>BMC Medical Genomics</i> , <b>2017</b> , 10, 48	3.7	19
35	Federated Tensor Factorization for Computational Phenotyping. <i>KDD: Proceedings</i> , <b>2017</b> , 2017, 887-895	6.8	30
34	SCOTCH: Secure Counting Of encryptEd genomIc data using a Hybrid approach <b>2017</b> , 2017, 1744-1753	0.7	1
33	Secure Multi-pArty Computation Grid LOGistic REgression (SMAC-GLORE). <i>BMC Medical Informatics and Decision Making</i> , <b>2016</b> , 16 Suppl 3, 89	3.6	27
32	HEALER: homomorphic computation of EXAct Logistic rEgRession for secure rare disease variants analysis in GWAS. <i>Bioinformatics</i> , <b>2016</b> , 32, 211-8	7.2	37
31	Lessons Learned for Online Health Community Moderator Roles: A Mixed-Methods Study of Moderators Resigning From WebMD Communities. <i>Journal of Medical Internet Research</i> , <b>2016</b> , 18, e247	7.6	25
30	A Predictive Model for Medical Events Based on Contextual Embedding of Temporal Sequences. <i>JMIR Medical Informatics</i> , <b>2016</b> , 4, e39	3.6	26
29	Protecting genomic data analytics in the cloud: state of the art and opportunities. <i>BMC Medical Genomics</i> , <b>2016</b> , 9, 63	3.7	30
28	Grid multi-category response logistic models. <i>BMC Medical Informatics and Decision Making</i> , <b>2015</b> , 15, 10	3.6	11
27	WebDISCO: a web service for distributed cox model learning without patient-level data sharing. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2015</b> , 22, 1212-9	8.6	58
26	Privacy-preserving GWAS analysis on Federated genomic datasets. <i>BMC Medical Informatics and Decision Making</i> , <b>2015</b> , 15 Suppl 5, S2	3.6	30
25	FORESEE: Fully Outsourced secuRe gENome Study basEd on homomorphic Encryption. <i>BMC Medical Informatics and Decision Making</i> , <b>2015</b> , 15 Suppl 5, S5	3.6	32
24	Differentially Private Histogram Publication For Dynamic Datasets: An Adaptive Sampling Approach <b>2015</b> , 2015, 1001-1010	4.5	23
23	pSCANNER: patient-centered Scalable National Network for Effectiveness Research. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2014</b> , 21, 621-6	8.6	59
22	DPSynthesizer: Differentially Private Data Synthesizer for Privacy Preserving Data Sharing. <i>Proceedings of the VLDB Endowment</i> , <b>2014</b> , 7, 1677-1680	3.1	24
21	A community assessment of privacy preserving techniques for human genomes. <i>BMC Medical Informatics and Decision Making</i> , <b>2014</b> , 14 Suppl 1, S1	3.6	37
20	Privacy preserving RBF kernel support vector machine. <i>BioMed Research International</i> , <b>2014</b> , 2014, 82737	3.1	14

19	Differentially Private Synthesization of Multi-Dimensional Data using Copula Functions <b>2014</b> , 2014, 475-486	11
18	Expectation Propagation LOGistic REgression (EXPLORER): distributed privacy-preserving online model learning. <i>Journal of Biomedical Informatics</i> , <b>2013</b> , 46, 480-96	10.2 47
17	SHARE: system design and case studies for statistical health information release. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2013</b> , 20, 109-16	8.6 26
16	WebGLORE: a web service for Grid LOGistic REgression. <i>Bioinformatics</i> , <b>2013</b> , 29, 3238-40	7.2 30
15	Privacy-preserving heterogeneous health data sharing. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2013</b> , 20, 462-9	8.6 30
14	Structured Set Intra Prediction With Discriminative Learning in a Max-Margin Markov Network for High Efficiency Video Coding. <i>IEEE Transactions on Circuits and Systems for Video Technology</i> , <b>2013</b> , 23, 1941-1956	6.4 2
13	Privacy technology to support data sharing for comparative effectiveness research: a systematic review. <i>Medical Care</i> , <b>2013</b> , 51, S58-65	3.1 26
12	Identifying inference attacks against healthcare data repositories. <i>AMIA Summits on Translational Science Proceedings</i> , <b>2013</b> , 2013, 262-6	1.1 8
11	Differential-Private Data Publishing Through Component Analysis <b>2013</b> , 6, 19-34	7
10	Genomes in the cloud: balancing privacy rights and the public good. <i>AMIA Summits on Translational Science Proceedings</i> , <b>2013</b> , 2013, 128	1.1 3
9	Calibrating predictive model estimates to support personalized medicine. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2012</b> , 19, 263-74	8.6 52
8	iDASH: integrating data for analysis, anonymization, and sharing. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2012</b> , 19, 196-201	8.6 114
7	A patient-driven adaptive prediction technique to improve personalized risk estimation for clinical decision support. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2012</b> , 19, e137-44	8.6 12
6	Grid Binary LOGistic REgression (GLORE): building shared models without sharing data. <i>Journal of the American Medical Informatics Association: JAMIA</i> , <b>2012</b> , 19, 758-64	8.6 107
5	A collaborative framework for Distributed Privacy-Preserving Support Vector Machine learning <b>2012</b> , 2012, 1350-9	0.7 5
4	Privacy-preserving SVM using nonlinear kernels on horizontally partitioned data <b>2006</b> ,	109
3	Privacy-Preserving SVM Classification on Vertically Partitioned Data. <i>Lecture Notes in Computer Science</i> , <b>2006</b> , 647-656	0.9 96
2	Secure Logistic Regression Based on Homomorphic Encryption: Design and Evaluation (Preprint)	6

1      Deep Learning for Alzheimer’s Disease Drug Repurposing using Knowledge Graph and Multi-level Evidence      1