

# Xin Gao

## List of Publications by Year in descending order

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36  
papers

601  
citations

840776

11  
h-index

642732

23  
g-index

37  
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37  
docs citations

37  
times ranked

580  
citing authors

#	ARTICLE	IF	CITATIONS
1	An Empirical Comparative Assessment of Inter-Rater Agreement of Binary Outcomes and Multiple Raters. <i>Symmetry</i> , 2022, 14, 262.	2.2	11
2	Graphical-model based high dimensional generalized linear models. <i>Electronic Journal of Statistics</i> , 2021, 15, .	0.7	1
3	Composite likelihood ratio testing under nonstandard conditions using tangent cones. <i>Stat</i> , 2021, 10, e375.	0.4	2
4	Penalized composite likelihood for colored graphical Gaussian models. <i>Statistical Analysis and Data Mining</i> , 2021, 14, 366-378.	2.8	2
5	On the Admissibility of Simultaneous Bootstrap Confidence Intervals. <i>Symmetry</i> , 2021, 13, 1212.	2.2	1
6	An Online Risk Tool for Predicting Type 2 Diabetes Mellitus. <i>International Journal of Diabetology</i> , 2021, 2, 123-129.	2.0	1
7	Media coverage and investment efficiency. <i>Journal of Empirical Finance</i> , 2021, 63, 270-293.	1.8	46
8	A Generalized Linear Mixed Model Approach to Assess Emerald Ash Borer Diffusion. <i>ISPRS International Journal of Geo-Information</i> , 2020, 9, 414.	2.9	2
9	Multivariate Mixed Response Model with Pairwise Composite-Likelihood Method. <i>Stats</i> , 2020, 3, 203-220.	0.9	6
10	Bayesian model selection approach for coloured graphical Gaussian models. <i>Journal of Statistical Computation and Simulation</i> , 2020, 90, 2631-2654.	1.2	3
11	On the application of a novel linear mixture model on laser-induced breakdown spectroscopy: Implications for Mars. <i>Journal of Chemometrics</i> , 2019, 33, e3174.	1.3	10
12	Predictive models for diabetes mellitus using machine learning techniques. <i>BMC Endocrine Disorders</i> , 2019, 19, 101.	2.2	148
13	FusionLearn: a biomarker selection algorithm on cross-platform data. <i>Bioinformatics</i> , 2019, 35, 4465-4468.	4.1	3
14	Composite likelihood model comparison test under fixed and local alternatives. <i>Stat</i> , 2018, 7, e182.	0.4	0
15	Approximate Bayesian estimation in large coloured graphical Gaussian models. <i>Canadian Journal of Statistics</i> , 2018, 46, 176-203.	0.9	0
16	Mixture Markov regression model with application to mosquito surveillance data analysis. <i>Biometrical Journal</i> , 2017, 59, 462-477.	1.0	3
17	Data integration with high dimensionality. <i>Biometrika</i> , 2017, 104, 251-272.	2.4	21
18	Multiple Comparisons Using Composite Likelihood in Clustered Data. <i>International Journal of Biostatistics</i> , 2016, 12, .	0.7	0

#	ARTICLE	IF	CITATIONS
19	Statistical Method for Integrative Platform Analysis: Application to Integration of Proteomic and Microarray Data. <i>Methods in Molecular Biology</i> , 2016, 1362, 199-207.	0.9	2
20	Estimation of Symmetry-Constrained Gaussian Graphical Models: Application to Clustered Dense Networks. <i>Journal of Computational and Graphical Statistics</i> , 2015, 24, 909-929.	1.7	9
21	Pilot Testing of a Psycho-educational Telephone Intervention for Women Receiving Uninformative BRCA1/2 Genetic Test Results. <i>Canadian Journal of Nursing Research</i> , 2015, 47, 53-71.	1.5	4
22	Simultaneous model selection and estimation for mean and association structures with clustered binary data. <i>Stat</i> , 2013, 2, 102-118.	0.4	1
23	Multiple-platform data integration method with application to combined analysis of microarray and proteomic data. <i>BMC Bioinformatics</i> , 2012, 13, 320.	2.6	14
24	Composite Likelihood Bayesian Information Criteria for Model Selection in High-Dimensional Data. <i>Journal of the American Statistical Association</i> , 2010, 105, 1531-1540.	3.1	91
25	Transition Dependency: A Gene-Gene Interaction Measure for Times Series Microarray Data. <i>Eurasip Journal on Bioinformatics and Systems Biology</i> , 2009, 2009, 1-12.	1.4	4
26	Asymptotic Behavior of Tail Density for Sum of Correlated Lognormal Variables. <i>International Journal of Mathematics and Mathematical Sciences</i> , 2009, 2009, 1-28.	0.7	29
27	Application of model selection technique in chemogenomic data analysis. <i>Statistical Analysis and Data Mining</i> , 2009, 2, 186-191.	2.8	0
28	Nonparametric multiple comparison procedures for unbalanced one-way factorial designs. <i>Journal of Statistical Planning and Inference</i> , 2008, 138, 2574-2591.	0.6	77
29	Nonparametric multiple comparison procedures for unbalanced two-way layouts. <i>Journal of Statistical Planning and Inference</i> , 2008, 138, 3674-3686.	0.6	21
30	A Nonparametric Procedure for the Two-Factor Mixed Model with Missing Data. <i>Biometrical Journal</i> , 2007, 49, 774-788.	1.0	7
31	Efficient Evaluation of BER for Arbitrary Modulation and Signaling in Fading Channels. <i>IEEE Transactions on Communications</i> , 2007, 55, 2061-2064.	7.8	11
32	CTH12-1: Efficient Calculation of BER for Arbitrary Modulation and Signaling in Fading Channels. <i>IEEE Global Telecommunications Conference (GLOBECOM)</i> , 2006, , .	0.0	0
33	Construction of null statistics in permutation-based multiple testing for multi-factorial microarray experiments. <i>Bioinformatics</i> , 2006, 22, 1486-1494.	4.1	16
34	Nonparametric tests for differential gene expression and interaction effects in multi-factorial microarray experiments. <i>BMC Bioinformatics</i> , 2005, 6, 186.	2.6	13
35	A nonparametric test for interaction in two-way layouts. <i>Canadian Journal of Statistics</i> , 2005, 33, 529-543.	0.9	16
36	A Unified Nonparametric Approach for Unbalanced Factorial Designs. <i>Journal of the American Statistical Association</i> , 2005, 100, 926-941.	3.1	26