

# Gang Li

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4671457/publications.pdf>

Version: 2024-02-01

41  
papers

1,771  
citations

567144

15  
h-index

315616

38  
g-index

41  
all docs

41  
docs citations

41  
times ranked

2780  
citing authors

| #  | ARTICLE  | IF   | CITATIONS |
|----|--|------|-----------|
| 1  | Neoadjuvant anti-PD-1 immunotherapy promotes a survival benefit with intratumoral and systemic immune responses in recurrent glioblastoma. <i>Nature Medicine</i> , 2019, 25, 477-486.                       | 15.2 | 932       |
| 2  | A Joint Model for Longitudinal Measurements and Survival Data in the Presence of Multiple Failure Types. <i>Biometrics</i> , 2008, 64, 762-771.  | 0.8  | 135       |
| 3  | An approach to joint analysis of longitudinal measurements and competing risks failure time data. <i>Statistics in Medicine</i> , 2007, 26, 2813-2835.   | 0.8  | 80        |
| 4  | A Bayesian approach to joint analysis of longitudinal measurements and competing risks failure time data. <i>Statistics in Medicine</i> , 2009, 28, 1601-1619.   | 0.8  | 69        |
| 5  | Maximum Likelihood Estimation in a Semiparametric Logistic/Proportional-Hazards Mixture Model. <i>Scandinavian Journal of Statistics</i> , 2005, 32, 59-75.  | 0.9  | 60        |
| 6  | On nonparametric likelihood ratio estimation of survival probabilities for censored data. <i>Statistics and Probability Letters</i> , 1995, 25, 95-104.  | 0.4  | 59        |
| 7  | A general joint model for longitudinal measurements and competing risks survival data with heterogeneous random effects. <i>Lifetime Data Analysis</i> , 2011, 17, 80-100.                                   | 0.4  | 43        |
| 8  | Latent Subgroup Analysis of a Randomized Clinical Trial through a Semiparametric Accelerated Failure Time Mixture Model. <i>Biometrics</i> , 2013, 69, 52-61.  | 0.8  | 39        |
| 9  | Simultaneous Estimation and Variable Selection for Interval-Censored Data With Broken Adaptive Ridge Regression. <i>Journal of the American Statistical Association</i> , 2020, 115, 204-216.                | 1.8  | 38        |
| 10 | Joint Modeling of Longitudinal and Time-to-Event Data. , 0, , .  |      | 37        |
| 11 | Extreme learning machine Cox model for high-dimensional survival analysis. <i>Statistics in Medicine</i> , 2019, 38, 2139-2156.  | 0.8  | 31        |
| 12 | Broken adaptive ridge regression and its asymptotic properties. <i>Journal of Multivariate Analysis</i> , 2018, 168, 334-351.  | 0.5  | 29        |
| 13 | A Unified Approach to Nonparametric Comparison of Receiver Operating Characteristic Curves for Longitudinal and Clustered Data. <i>Journal of the American Statistical Association</i> , 2008, 103, 705-713. | 1.8  | 23        |
| 14 | Empirical likelihood analysis of the Buckley-James estimator. <i>Journal of Multivariate Analysis</i> , 2008, 99, 649-664.   | 0.5  | 22        |
| 15 | Efficient Regularized Regression with $L_1$ Penalty for Variable Selection and Network Construction. <i>Computational and Mathematical Methods in Medicine</i> , 2016, 2016, 1-11.                           | 0.7  | 22        |
| 16 | Longitudinal data analysis with non-ignorable missing data. <i>Statistical Methods in Medical Research</i> , 2016, 25, 205-220.  | 0.7  | 19        |
| 17 | Empirical Likelihood for Censored Linear Regression and Variable Selection. <i>Scandinavian Journal of Statistics</i> , 2015, 42, 798-812.   | 0.9  | 16        |
| 18 | A semiparametric linear transformation model to estimate causal effects for survival data. <i>Canadian Journal of Statistics</i> , 2014, 42, 18-35.  | 0.6  | 11        |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 19 | An oracle property of the Nadaraya-Watson kernel estimator for high-dimensional nonparametric regression. <i>Scandinavian Journal of Statistics</i> , 2019, 46, 735-764.   | 0.9 | 9         |
| 20 | Prediction Accuracy Measures for a Nonlinear Model and for Right-Censored Time-to-Event Data. <i>Journal of the American Statistical Association</i> , 2019, 114, 1815-1825.   | 1.8 | 9         |
| 21 | Scalable Algorithms for Large Competing Risks Data. <i>Journal of Computational and Graphical Statistics</i> , 2021, 30, 685-693.  | 0.9 | 9         |
| 22 | Comments on: A review on empirical likelihood methods for regression. <i>Test</i> , 2009, 18, 463-467.   | 0.7 | 8         |
| 23 | Analysis of two-sample censored data using a semiparametric mixture model. <i>Acta Mathematicae Applicatae Sinica</i> , 2009, 25, 389-398.   | 0.4 | 7         |
| 24 | Variable selection for recurrent event data with broken adaptive ridge regression. <i>Canadian Journal of Statistics</i> , 2018, 46, 416-428.  | 0.6 | 7         |
| 25 | A surrogate $\ell_1$ -sparse Cox's regression with applications to sparse high-dimensional massive sample size time-to-event data. <i>Statistics in Medicine</i> , 2020, 39, 675-686.                                  | 0.8 | 7         |
| 26 | An Empirical Likelihood Method for Semiparametric Linear Regression with Right Censored Data. <i>Computational and Mathematical Methods in Medicine</i> , 2013, 2013, 1-9.   | 0.7 | 6         |
| 27 | A new joint screening method for right-censored time-to-event data with ultra-high dimensional covariates. <i>Statistical Methods in Medical Research</i> , 2020, 29, 1499-1513.                                       | 0.7 | 6         |
| 28 | Joint Inference for Competing Risks Survival Data. <i>Journal of the American Statistical Association</i> , 2016, 111, 1289-1300.  | 1.8 | 5         |
| 29 | Two-step hypothesis testing to detect gene-environment interactions in a genome-wide scan with a survival endpoint. <i>Statistics in Medicine</i> , 2022, 41, 1644-1657.   | 0.8 | 5         |
| 30 | Broken adaptive ridge regression for right-censored survival data. <i>Annals of the Institute of Statistical Mathematics</i> , 2022, 74, 69-91.  | 0.5 | 4         |
| 31 | Non-parametric Estimation of a Survival Function with Two-stage Design Studies. <i>Scandinavian Journal of Statistics</i> , 2008, 35, 193-211.   | 0.9 | 3         |
| 32 | Nonparametric inference for assessing treatment efficacy in randomized clinical trials with a time-to-event outcome and all-or-none compliance. <i>Biometrika</i> , 2012, 99, 393-404.                                 | 1.3 | 3         |
| 33 | Sample size determination for jointly testing a cause-specific hazard and the all-cause hazard in the presence of competing risks. <i>Statistics in Medicine</i> , 2018, 37, 1389-1401.                                | 0.8 | 3         |
| 34 | Variable Selection in Threshold Regression Model with Applications to HIV Drug Adherence Data. <i>Statistics in Biosciences</i> , 2020, 12, 376-398.   | 0.6 | 3         |
| 35 | A scalable surrogate $\ell_1$ -sparse regression method for generalized linear models with applications to large scale data. <i>Journal of Statistical Planning and Inference</i> , 2021, 213, 262-281.                |     |           |
| 36 | A flexible joint model for multiple longitudinal biomarkers and a time-to-event outcome: With applications to dynamic prediction using highly correlated biomarkers. <i>Biometrical Journal</i> , 2021, 63, 1575-1586. | 0.6 | 3         |

| #  | ARTICLE  | IF  | CITATIONS |
|----|--|-----|-----------|
| 37 | Sure joint feature screening in nonparametric transformation model for right censored data. Canadian Journal of Statistics, 2021, 49, 549-565.   | 0.6 | 2         |
| 38 | Efficient Algorithms and Implementation of a Semiparametric Joint Model for Longitudinal and Competing Risk Data: With Applications to Massive Biobank Data. Computational and Mathematical Methods in Medicine, 2022, 2022, 1-12. | 0.7 | 2         |
| 39 | Simultaneous estimation and variable selection for incomplete event history studies. Journal of Multivariate Analysis, 2019, 171, 350-361.   | 0.5 | 1         |
| 40 | SEMIPARAMETRIC ADDITIVE RISKS REGRESSION FOR TWO-STAGE DESIGN SURVIVAL STUDIES. Statistica Sinica, 2010, 20, 1581-1607.  | 0.2 | 1         |
| 41 | A New $\hat{\alpha}_0$ -Regularized Log-Linear Poisson Graphical Model with Applications to RNA Sequencing Data. Journal of Computational Biology, 2021, 28, 880-891.  | 0.8 | 0         |