

Nan Xiao

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Academic degrees

2008 – 2012 Bachelor of Science (Statistics), Central South University. *Changsha, China*

Studies

2015 – 2016 Ph.D. Student (Human Genetics), University of Chicago. *Chicago, IL, U.S.*
2012 – 2015 Ph.D. Student (Statistics), Central South University. *Changsha, China*

Work experience

2016 – Genomic Data Scientist. Seven Bridges Genomics, Inc. *Cambridge, MA, U.S.*

Publications

Preprints

2016 **Nan Xiao**, Q.-S. Xu, and M.-Z. Li (2016). “hdnom: Building nomograms for penalized Cox models with high-dimensional survival data”. *bioRxiv*. doi: [10.1101/065524](https://doi.org/10.1101/065524).

Journal articles

2015 **Nan Xiao** and Q.-S. Xu (2015). “Multi-step adaptive elastic-net: reducing false positives in high-dimensional variable selection”. *Journal of Statistical Computation and Simulation*. doi: [10.1080/00949655.2015.1016944](https://doi.org/10.1080/00949655.2015.1016944).

2015 **Nan Xiao**, D.-S. Cao, M.-F. Zhu, and Q.-S. Xu (2015). “protr/ProtrWeb: R package and web server for generating various numerical representation schemes of protein sequence”. *Bioinformatics*. doi: [10.1093/bioinformatics/btv042](https://doi.org/10.1093/bioinformatics/btv042).

2014 D.-S. Cao*, **Nan Xiao***, Q.-S. Xu and A. F. Chen (2014). “Rcpi: R/Bioconductor package to generate various descriptors of proteins, compounds, and their interactions”, *Bioinformatics*. ***Joint first authors**. doi: [10.1093/bioinformatics/btu624](https://doi.org/10.1093/bioinformatics/btu624).

2015 D.-S. Cao, **Nan Xiao**, Y.-J. Li, W.-B. Zeng, Y.-Z. Liang, A.-P. Lu, Q.-S. Xu, A. F. Chen (2015). “Integrating multiple evidence sources to predict adverse drug reactions based on systems pharmacology model”. *CPT: Pharmacometrics & Systems Pharmacology*. doi: [10.1002/psp4.12002](https://doi.org/10.1002/psp4.12002).

2015 J.-B. Wang, D.-S. Cao, M.-F. Zhu, Y.-H. Yun, **Nan Xiao**, Y.-Z. Liang (2015). “*In silico* evaluation of logD_{7.4} and comparison with other prediction methods”. *Journal of Chemometrics*. doi: [10.1002/cem.2718](https://doi.org/10.1002/cem.2718).

- 2015 L. Shen, D.-S. Cao, Q.-S. Xu, X. Huang, **Nan Xiao**, Y.-Z. Liang (2015). “A novel local manifold-ranking based k -NN for modeling the regression between bioactivity and molecular descriptors”. *Chemometrics and Intelligent Laboratory Systems*. doi: [10.1016/j.chemolab.2015.12.005](https://doi.org/10.1016/j.chemolab.2015.12.005).

Book translations

- 2016 Max Kuhn and Kjell Johnson (2016). *Applied Predictive Modeling*. (Hui Lin, Yi-Xuan Qiu, En-Chi Ma, **Nan Xiao**, & Vivian Zhang, Trans.). China Machine Press (Original work published in 2013). ISBN: [978-7-1115-3342-9](https://www.isbn-international.org/product/978-7-1115-3342-9).
- 2014 Winston Chang (2014). *R Graphics Cookbook*. (**Nan Xiao**, Yi-Shuo Deng, Tai-Yun Wei, & Yi-Xuan Qiu, Trans.). Posts and Telecom Press (Original work published in 2013). ISBN: [978-7-115-34227-0](https://www.isbn-international.org/product/978-7-115-34227-0).
- 2013 Hadley Wickham (2013). *ggplot2: Elegant Graphics for Data Analysis*. (Tai-Yun Wei, Yi-Xuan Qiu, **Nan Xiao**, Tao Gao, & Wei-Cheng Zhu, Trans.). Xi'an Jiaotong University Press (Original work published in 2010). ISBN: [978-7-5605-4969-9](https://www.isbn-international.org/product/978-7-5605-4969-9).
- 2013 Robert Kabacoff (2013). *R in Action: Data Analysis and Graphics with R*. (Tao Gao, **Nan Xiao**, & Gang Chen, Trans.). Posts and Telecom Press (Original work published in 2011). ISBN: [978-7-115-29990-1](https://www.isbn-international.org/product/978-7-115-29990-1).

Software and web applications

R/Bioconductor packages

hdnom

- 2015 – R package building nomograms for high-dimensional survival data, with support for model validation, calibration, and comparison. Website: <http://hdnom.org>. Available from CRAN: <https://cran.r-project.org/package=hdnom>.

msaenet

- 2016 – R package implementing the multi-step adaptive elastic-net (MSAENet) algorithm for feature selection in high-dimensional regressions. Website: <http://msaenet.com>. Available from CRAN: <https://cran.r-project.org/package=msaenet>.

protr

- 2012 – R package generating various numerical representation schemes of protein sequence for bioinformatics and proteochemometrics research. Website: <http://nanx.me/protr/>. Available from CRAN: <https://cran.r-project.org/package=protr>.

Rcpi

- 2013 – R/Bioconductor package with comprehensive integration of bioinformatics and cheminformatics tools for drug discovery. Website: <http://nanx.me/Rcpi/>. Available from Bioconductor: <https://bioconductor.org/packages/Rcpi>.

- ggsci**
2016 – R package offering a collection of ggplot2 color palettes inspired by scientific journals, data visualization libraries, and science fiction TV shows. Website: <http://ggsci.net>. Available from CRAN: <https://cran.r-project.org/package=ggsci>.
- liftr**
2015 – R package for dockerizing R Markdown documents with support for Rabix in reproducible research. Website: <http://liftr.me>. Available from CRAN: <https://cran.r-project.org/package=liftr>.
- enpls**
2014 – R package for ensemble partial least squares regression, a unified framework for feature selection, outlier detection, and ensemble learning. Website: <http://enpls.org>. Available from CRAN: <https://cran.r-project.org/package=enpls>.
- sevenbridges**
2016 – Seven Bridges API client, CWL schema, Meta schema, and SDK helper in R. Available from Bioconductor: <https://bioconductor.org/packages/sevenbridges>.
- sbgr**
2015 – R Client for Seven Bridges Genomics API (v1). Available from GitHub: <https://github.com/road2stat/sbgr>.
- grex**
2016 – R package for mapping Ensembl gene IDs in Genotype-Tissue Expression (GTEx) data to Entrez IDs, HGNC symbols, and UniProt IDs. Website: <http://nanx.me/grex/>. Available from CRAN: <https://cran.r-project.org/package=grex>.
- RECA**
2012 – R package for relevant component analysis (RCA) in supervised distance metric learning and supervised dimensionality reduction. Available from CRAN: <https://cran.r-project.org/package=RECA>.

Web applications

- hdnom.io**
2015 – Shiny web application for building penalized Cox models for high-dimensional data with survival outcomes. Streamlined the process of nomogram building, model validation, calibration, and comparison. Available at: <http://hdnom.io>.
- ProtrWeb**
2013 – Shiny web application for computing 20 types of protein sequence-derived structural and physicochemical features in bioinformatics. Available at <http://protr.org>.
- TargetNet**
2014 – Shiny web application for drug target identification with large-scale public binding affinities data. Available at <http://targetnet.org>.

ImgSVD

- 2014 – Shiny web application for image compression via singular value decomposition.
Collaboration with *Yihui Xie*, *Yixuan Qiu*, and *Tong He*.
Available at <https://yihui.shinyapps.io/imgsvd>.

Signify

- 2015 Shiny web application for making your (>0.05) p -values sound significant.
Available at <https://nanx.shinyapps.io/signify>.

Professional services

- 2016 – **Reviewer.** *The R Journal*
2015 – **Reviewer.** *Journal of Statistical Computation and Simulation*
2014 **Organizing Committee Member.** *The 7th China R Conference (Beijing)*
2013 **Co-Organizer.** *The 6th China R Conference (Beijing)*

Conference & workshop presentations

- 2016 **hdnom.io: High-Dimensional Survival Modeling with Shiny.**
Lightning talk. Shiny Developer Conference, Stanford University. January 2016.
2015 **Introduction to Reproducible Research in Bioinformatics.**
Invited talk. 2015 Bioinformatics Workshop, Center for Research Informatics,
University of Chicago. December 3, 2015.
2013 **Web Scraping with R**
The 6th China R Conference, Renmin University of China, Beijing, China. May 2013.

Honors and awards

- 2013 National scholarship for graduate students (5%).
Highest award the government established for graduate students in China.
2013 Outstanding graduate student award (5%). Central South University, China.
2011 1st Prize (top 5%) in Hunan Contest District in China Undergraduate Mathematical
Contest in Modeling (CUMCM). With *Guan-Chun Liu* and *Tian-Yu Zhao*.
2010 Meritorious award (2nd in 1,000+) of Central China Undergraduate Mathematical
Contest in Modeling (CCUMCM). With *Tao Gao* and *Chen-Xi Guo*.

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