

Nan Xiao

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Education

2015 –	Ph.D. student (Human Genetics), University of Chicago.	Chicago, IL, U.S.
2012 – 2015	Master student (Statistics), Central South University.	Changsha, China
2008 – 2012	Bachelor of Science (Statistics), Central South University.	Changsha, China

Work experience

2015	Intern. Seven Bridges Genomics, Inc. Advisor: Tengfei Yin.	Cambridge, MA, U.S.
	<ul style="list-style-type: none">• Developed R/Bioconductor package <i>sbgr</i>: R client for SBG Platform / NCI Cancer Genomics Cloud Platform API; the first shipped official language binding for SBG API.• Developed R package <i>liftr</i>: dockerizing R Markdown documents with support for Rabix. First enabled OS-level reproducibility for bioinformatics & statistical data analysis.• Presented the packages at the BioC 2015 conference (workshop session and poster session), Fred Hutchinson Cancer Research Center, Seattle, WA.	

Publications

Journal articles

2015	Nan Xiao and Q.-S. Xu (2015). “Multi-step adaptive elastic-net: reducing false positives in high-dimensional variable selection”. <i>Journal of Statistical Computation and Simulation</i> . doi: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”. <i>Bioinformatics</i> . doi: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”, <i>Bioinformatics</i> . *Joint first authors. [Package downloaded over 6,000 times.] doi: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”. <i>CPT: Pharmacometrics & Systems Pharmacology</i> . doi: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”. <i>Journal of Chemometrics</i> . doi:10.1002/cem.2718	

Book translations

- 2015 Max Kuhn and Kjell Johnson (2015). *Applied Predictive Modeling*. (En-Chi Ma, Yi-Xuan Qiu, **Nan Xiao**, Hui Lin, & Vivian Zhang, Trans.). China Machine Press (Original work published in 2013). In press.
- 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).
- 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).
- 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). [Sold 27,500 copies.]

Conference presentations & posters

- 2015 **Introduction to Reproducible Research in Bioinformatics**.
Invited talk. 2015 Bioinformatics Workshop, Center for Research Informatics, University of Chicago. December 3, 2015.
- 2015 **Nan Xiao and Tengfei Yin. liftr: Reproducible Bioinformatics and Statistical Data Analysis with Docker, Rabix, and knitr** (Poster).
BioC 2015, Fred Hutchinson Cancer Research Center, Seattle, WA. July 2015.
- 2015 **liftr & sbgr kickstart** (Workshop with Dan Tenenbaum and Tengfei Yin).
BioC 2015, Fred Hutchinson Cancer Research Center, Seattle, WA. July 2015.
- 2014 **Integrated Pipeline for Systems Pharmacology in R/Bioconductor**
The 7th China R Conference, Renmin University of China, Beijing, China. May 2014.
- 2013 **Web Scraping with R**
The 6th China R Conference, Renmin University of China, Beijing, China. May 2013.
- 2009 **Security Issues and Vulnerabilities of the R Environment**
The 2nd China R Conference, Renmin University of China, Beijing & East China Normal University, Shanghai. December 2009.

Software and web applications

R/Bioconductor packages

hdnom

- 2015 – R package building nomograms for high-dimensional Cox models, with support for model validation and calibration. Project website: <http://hdnom.org>.
Availbale from CRAN: <https://cran.r-project.org/web/packages/hdnom/>.

- liftr**
2015 – R package for dockerizing R Markdown documents with support for Rabin in reproducible research. Project website: <http://liftr.me>.
Available from CRAN: <https://cran.r-project.org/web/packages/liftr/>.
- sbgr**
2015 – R Client for Seven Bridges Genomics API.
Available from Bioconductor: <http://bioconductor.org/packages/sbgr/> and
GitHub: <https://github.com/road2stat/sbgr>.
- Rcpi**
2013 – 2014 R/Bioconductor package as a comprehensive molecular informatics platform for systems pharmacology and drug discovery. [Downloaded over 6,000 times by February 2015.]
Available from Bioconductor: <http://bioconductor.org/packages/Rcpi>.
- protr**
2012 – 2014 R package generating various numerical representation schemes of protein sequence for bioinformatics and proteochemometrics research.
Available from CRAN: <http://cran.r-project.org/web/packages/protr/>.
- enpls**
2014 R package for ensemble partial least squares regression, a unified framework for feature selection, outlier detection, and ensemble learning.
Available from CRAN: <http://cran.r-project.org/web/packages/enpls/>.
- RECA**
2012 – 2014 R package for relevant component analysis (RCA) in supervised distance metric learning and supervised dimensionality reduction.
Available from CRAN: <http://cran.r-project.org/web/packages/RECA/>.

Web applications

- hdnom.io**
2015 Web application for building penalized Cox models for high-dimensional data with survival outcomes. The web app streamlined the process of nomogram building, model validation, model calibration, and reproducible report generation.
Available at: <http://hdnom.io>.
Included by Shiny user showcase: <https://rstudio.com/products/shiny/shiny-user-showcase/>
- ProtrWeb**
2013 – 2014 Web application for computing 20 types of protein sequence-derived structural and physicochemical features in bioinformatics. [Over 2,000 users from 40 countries.]
Available at <http://protrweb.scbdd.com>.
- TargetNet**
2014 Web application for predicting the binding of 623 drug targets for given molecules. Driven by large-scale machine learning modeling of chemogenomics data.
Available at: <http://targetnet.scbdd.com>.

ImgSVD

2014 Web application for image compression via singular value decomposition. Collaborated with *Yihui Xie* (RStudio Inc.), *Yixuan Qiu* (Purdue University), and *Tong He* (Simon Fraser University). Available at <https://yihui.shinyapps.io/imgsvd/>.

Honors and awards

2014 Best poster award. *Genes and associated environmental factors affecting skin aging in different ethnic groups*. CAS-MPG Partner Institute for Computational Biology Scientific Exchange Meeting, Shanghai. With *Miaozhu Li*.

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, Changsha, 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*.

Professional services

2015 – **Reviewer.** *Journal of Statistical Computation and Simulation*

2014 **Contributing Editor.** *ICSA (International Chinese Statistical Association) Bulletin*

2014 **Organizing Committee Member.** *The 7th China R Conference (Beijing)*. <http://china-r.org>

2013 **Co-Organizer.** *The 6th China R Conference (Beijing)*

2010 – **Editor.** *Capital of Statistics*, <http://cos.name>. Largest online community on statistics in China.

2009 **Volunteer.** *The 2nd China R Conference (Beijing, Shanghai)*

Professional skills

Research	<ul style="list-style-type: none">• High-dimensional linear models• High-performance statistical computing	<ul style="list-style-type: none">• Statistical machine learning methods• Statistical graphics and data visualization
Programming	<ul style="list-style-type: none">• R/Bioconductor package development (6 years)• Complex web application development with R/Shiny (3 years)• Good knowledge of Docker, Python, PostgreSQL, T_EX and shell programming (4 years)• Experience of Linux server setup, management and application deployment (4 years)	

Last revision: December, 2015