WENCAN ZHU

Ph.D. in Applied Mathematics

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RESEARCH EXPERIENCE

University of Paris-Saclay & Sanofi R&D PhD Research

- **2019 Present**
- Paris, France
- Dissertation: development of novel machine learning methods for biomarker selection in high-dimensional settings with strong correlations.
 - WLasso and Generalized Elastic Net: select progsnotic biomarkers by whitening the design matrix (linear regression model).
 - PPLasso: simultaneously select prognostic and predictive biomarkers (ANCOVA-type model).
 - WLogit: extension of WLasso to binary classification.
- Proiec
 - Applied the methods above to projects of Sanofi (RNAseq and Olink data as examples)
- Supervision
 - supervised a graduate internship on the comparison of R package LIMMA and classical ANCOVA model.

Sanofi R&D

Biostatistician

- **2018 2019**
- Chilly-Mazarin, France
- Statistical modeling with Omics data (RNAseq, GWAS):
 - Preprocessing
 - Identification of associated biomarkers
 - Machine learning regression/classification

ENSAI & Sanofi R&D

Graduate internship

2018

- Chilly-Mazarin, France
- Comparison of penalized regression and screening approaches for the identification of prognostic biomarkers
 - Compared and studied performance of different methods (Lasso, SCAD, Elastic-Net, Adaptive Ridge, Adaptive Lasso, univariate analysis) regarding the efficiency of selection and prediction.
 - · Investigated on pres-screening by SIS

INSEE (National Institute of Statistics and Economic Studies)

Summer internship

2017

- Lyon, France
- Worked in the methodology department to define strategies for statistical analyse and model development in spatial analysis.
- Analyzed data from census and surveys for prediction, by using mixed model with spatial correlations.
- Tried different generalized models: Poisson, Zero-inflated model.

EDUCATION

Ph.D. in Mathematics University of Paris-Saclay

= 2022

Paris, France

M.Sc. in Biostatistics

ENSAI (National School for Statistics and Data Analyse)

2018

Rennes, France

M.Sc. in Bioinformatics

Faculty of Medicine, Rennes 1 University

= 2018

Rennes, France

M.Sc. in Mathematics

Department of Mathematics, Tongji University

= 2018

Shanghai, China

B.Sc. (major) in Mathematics Department of Mathematics, Tongji University

= 2015

Shanghai, China

B.A. (minor) in Japanese

Department of Literature, Tongji University

= 2015

Shanghai, China

IT SKILLS

R Python

JAVA

C++

R Shiny

ATEX SAS LINUX

LANGUAGES

Mandarin English French Japanese German



PUBLICATIONS

Journal Articles

- Zhu, W., Lévy-Leduc, C., & Ternès, N. (2021). A variable selection approach for highly correlated predictors in high-dimensional genomic data. *Bioinformatics*, 37(16), 2238–2244.
- Zhu, W., Adjakossa, E., Lévy-Leduc, C., & Ternès, N. (2021). Sign consistency of the generalized elastic net estimator. Undergoing peer review. (arXiv: 2106.05454 [math.ST])
- Zhu, W., Lévy-Leduc, C., & Ternès, N. (2022). Identification of prognostic and predictive biomarkers in high-dimensional data with PPLasso. Undergoing peer review. (arXiv: 2202.01970 [stat.ME])
- Zhu, W., Lévy-Leduc, C., & Ternès, N. (2022). Variable selection in high-dimensional logistic regression models using a whitening approach. Submitted. (arXiv: 2206.14850 [stat.ME])
- Liu, F., Zhu, W., Shoaito, H., Chissey, A., Degrelle, S. A., & Fournier, T. (2020). Mining of combined human placental gene expression data across pregnancy, applied to PPAR signaling pathway. *Placenta*, 99, 157–165.
- Liu, F., Rouault, C., Guesnon, M., Zhu, W., Clément, K., Degrelle, S. A., & Fournier, T. (2020). Comparative Study of PPAR Targets in Human Extravillous and Villous Cytotrophoblasts. PPAR research, 9210748.
- Liu, F., Simasotchi, C., Vibert, F., Zhu, W., Gil, S., Degrelle, S. A., & Fournier, T. (2021). Age and Sex-Related Changes in Human First-Trimester Placenta Transcriptome and Insights into Adaptative Responses to Increased Oxygen. *International journal of molecular sciences*, 22(6), 2901.
- Liu, F., Rouault, C., Clément, K., Zhu, W., Degrelle, S. A., Charles, M. A., Heude, B., & Fournier, T. (2021). C1431T Variant of PPAR Is Associated with Preeclampsia in Pregnant Women. *Life (Basel, Switzerland)*, 11(10), 1052.

■ Book

• Zhu, W., Ardilly, P., Bouche, P (2018). Handbook of spatial analysis (Chapter 12. Small areas and spatial correlation). INSEE-Eurostat.

Package R

- **Zhu, W.** (2022). WLogit: Whitening logistic regression with regularization.
- **Zhu**, **W.**, Lévy-Leduc, C., & Ternès, N. (2022b). *PPLasso: Prognostic and predictive biomarker identification*.
- Zhu, W., Lévy-Leduc, C., & Ternès, N. (2020). WLasso: Whitening lasso for variable selection.

Conference Communications (oral presentation)

- ISCB (International Society for Clinical Biostatistics) Zhu, W., Lévy-Leduc, C., & Ternès, N. (2022). Identification of prognostic and predictive biomarkers in high-dimensional data with pplasso.
- JDS (Journées de Statistique) Zhu, W., Lévy-Leduc, C., & Ternès, N. (2022). Identification of prognostic and predictive biomarkers in high-dimensional data with pplasso.
- SMPGD (Statistical Methods for Post Genomic Data) Zhu, W., Lévy-Leduc, C., & Ternès, N. (2022). Wlasso: Variable selection for highly correlated predictors (application to genomic data).
- ISCB (International Society for Clinical Biostatistics) Zhu, W., Lévy-Leduc, C., & Ternès, N. (2020). A variable selection approach for highly correlated predictors in high-dimensional genomic data.

REFEREES

Céline Lévy-Leduc (Professor)

- @ AgroParisTech, INRAE, University of Paris-Saclay

Nils Ternès (Statistical Biomarker Leader)

- @ Sanofi R&D
 -] nils.ternes@sanofi.com

Pascal Ardilly (Director)

- @ Department of methodology, INSEE
 - □] pascal.ardilly@insee.fr

FELLOWSHIP

 Eiffel scholarship program of excellence, French Ministry for Europe and Foreign Affairs, 2016-2018