Liguang Zhu

Data Scientist

Liguang (Leon) Zhu
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Highlights

- Strong skills in analytics and solution prototyping with R, Python and Weka.

- Extensive experience in cross-disciplinary research, analytics and predictive modeling.

- Experienced in AWS, Microsoft Azure, high performance computing, and grid computing.

- Experienced in SQL, NoSQL, data warehousing.

- Exposed to Hadoop and its ecosystem

Experience

Mar 2010 - Present, Clatyon School of IT, Monash University
Researcher & PhD

Specialise in data science disciplines including machine learning, probabilistic modeling, predictive modeling and graph theory.

- Played a key role in researching and protyping of a novel feature engineering algorithm: Subsumption Resolution.

- Independently researched various predictive models on highly imbalanced biological data and implemented a tool for enzyme specificity prediction.

- Leading the design and development of a novel protein design pipeline by study high dimensional biological data.

- Designed one of the most stable protein molecules: FN3con.

- Designed a pharmaceutically promising molecule that challenges the current understanding of structure-folding relationship of proteins.

- Coordinated with research groups in different disciplines such as data science, structural biology, protein production unit, and chemical engineering.

Nov 2016 - Mar 2017, South East Water
Data Scientist Intern

Responsible for exploring statistics and machine learning solutions to risk prediction and budget modeling of the gravity sewerage system.

- Utilised customer retention analysis for system failure modeling.

- Utilised customer segmentation analysis to reduce selection-bias.

- Explored and analysed datasets with various machine learning techniques such as classification, clustering, association rule discovery.

- Sourced multiple datasets of CCTV inspections, historical renewal programs and system failure records from data warehouses.

- Designed and implemented a budgeting optimisation software.

- Utilised tableau for visual analytics and reporting.

Sep 2012 - Present, Monash University
Teaching Associate

Responsible for delivering, facilitating and managing tutorials, discussion, and practical experiments.

- Advanced algorithms and data structures
- Business intelligence and data warehousing
- Introduction to data science
- Introduction to computer science
- programming foundations in Python

Sep 2010 - Dec 2010, School of Biological Science, Monash University
Research Assistant

Responsible for the development and deployment of a portable database system on customised data collection devices for ecology research.

Sep 2010 - Dec 2012, Fleet Software & Services Pty Ltd
Freelancer Data Science Consultant

Involved in the design and development of a value prediction system based on historical fleet exchange data.

- Managed data cleaning and predictive modeling.
- Minimised the average prediction error within $600.

Education

2012 - present, Monash University
Doctorate of Philosophy

2008 - 2010, Monash University
Master of Computer Science (Minor Thesis in Data Science)

2003 - 2007, Beijing Jiaotong University, China
Bachelor of Engineering in Computer Science with Honours

References

Provide upon request.

Publications

- B. Porebski, S. Keleher, J. Hollins, A. Nickson, **L. Zhu**, et. al. (2016) *Smoothing a rugged protein folding landscape by sequence-based redesign.* Scientific Report, 6, Art. no. 33958.

- B.T. Porebski, A.A. Nickson, D.E. Hoke, M.R. Hunter, **L. Zhu**, S. McGowan, G.I. Webb, and A.M. Buckle. (2015). *Structural and dynamic properties that govern the stability of an engineered fibronectin type III domain.* Protein Engineering, Design and Selection. 28(3): 67-78. Oxford University Press.

- **L. Zhu**, B.T. Porebski, A.M. Buckle, G.I. Webb. (2013). *A probabilistic approach to In Silico protein design.* QMB E3: Enzyme Engineering and Evolution. Queenstown, New Zealand.

- B.T. Porebski, **L. Zhu**, D.E. Hoke, W. Dai, S. Keleher, N.A. Borg, S.P. Bottomley, G.I.Webb, A.M. Buckle. (2013). *A structural, biophysical and computational investigation of two sequence-based protein engineering methods.* Poster session presented at: The 38th Lorne Conference on Protein Structure and Function. Lorne, Victoria.

- F. Zheng, G.I. Webb, P. Suraweera, and **L. Zhu**. (2012). *Subsumption Resolution: An Efficient and Effective Technique for Semi-Naive Bayesian Learning.* Machine Learning 87(1): 93-125. Springer Netherlands.