

# Danlu Liu

GitHub: <https://github.com/liudanlu001> • LinkedIn: [www.linkedin.com/in/danluliu1993](http://www.linkedin.com/in/danluliu1993)

Tel: 573-823-6626 • Email: [dltb9@mail.missouri.edu](mailto:dltb9@mail.missouri.edu)

## OBJECTIVE

Enthusiastic data science research seeking for an internship data scientist position in 2020 to support business growth and success by leveraging my 5 years of experience in data analytics and machine learning.

## EDUCATION

<b>Ph.D. student</b> in Computer Science, University of Missouri, Columbia, MO (GPA 3.98/4.0)	08/2017-12/2021
<b>M.S.</b> in Computer Science, University of Missouri, Columbia, MO (GPA 4.0/4.0)	08/2015-05/2017
<b>B.S.</b> in Applied Mathematics, China University of Petroleum, Tsingtao, China	09/2011-06/2015

## EXPERIENCE

- **Graduate Research Assistant**, University of Missouri, Columbia, MO
  - 1) Exploratory data mining for subgroup cohort discoveries and decision support on EHR/EMR dataset.
    - Developed a novel subgroup discovery method slice and dice thousands of potential subpopulations and prioritize potential cohorts.
    - Recommended hundreds co-occurring risk factors patterns of each subgroup and provide interventionable insights.
    - Applications:
      - SFARI Autism data: 6 novel subgroups discovered among 2,591 probands' data with 15 phenotypes and 10000 genotypes.
      - Cerner HealthFacts glaucoma data: a list of explainable patterns discovered for glaucoma prediction among 131,257 patients' data with at least 12-year longitudinal history.
      - T1D Exchange Clinic Registry data (2010-12; 2015-17): Multiple risk factors discovered between paternal T1D and maternal T1D.
      - The Cancer Genome Atlas (TCGA) data: Significant risk factors discovered among 625 breast cancer RNA-Seq data.
  - 2) Build a novel indexing structure for risk pattern mining, which is 100× faster compared to the state of the art.
    - Performed gene expression analysis between 555 cancer patients and 100 normal breast tissues.
    - Identified 10 significant single/combined risk factors between Triple-negative and Luminal A cancer types.
  - 3) Conducted BigData ecosystem processing and analysis with large-scale GeoSpatial data
    - Building a large database for 300+ million data points all over United States by using HBase and Hadoop.
    - Performed large-scaled geospatial calculation by using R and Spark.
- **Graduate Teaching Assistant**, University of Missouri, Columbia, MO
  - 1) Assisted with CS4050-Design and Analysis of Algorithms course. (50+ students/semester) 05/2017-12/2019
  - 2) Assisted with CS4410-Theory of Computation course. (40+ students/semester) 01/2016-12/2019

## SKILLS

Programming: Scala, Python, C/C++, Java, R

Tools: Spark, Hadoop, HBase, Hive, RStudio, Jupyter, MATLAB, Tableau, Git, Linux

Statistical Skills: Multivariate Analysis, Generalized Linear Model (GLM), Time Series, Bayesian Analysis, Machine Learning, Association Rule Mining, Natural Language Processing (NLP), Recommender System

## PUBLICATIONS (2 MORE ON LINKEDIN)

**D. Liu**, W. Baskett, D. Beversdorf and C. Shyu, "Exploratory Data Mining for Subgroup Cohort Discoveries and Prioritization," in IEEE Journal of Biomedical and Health Informatics. (IF 2018: 4.217).

M. Raju, **D. Liu**, F. W. Fraunfelder and C. Shyu, "Discovering multifactorial associations with the development of age-related cataract using contrast mining," 2017 IEEE International Conference on Bioinformatics and Biomedicine (BIBM), Kansas City, MO, 2017, pp. 2297-2299.