Xi (Bill) Chen | Resume

L +1 857 209 1002 • \Box +1 309 455 4260 • \boxtimes billchenxi@gmail.com **O** https://billchenxi.github.io • https://www.linkedin.com/in/billchenxi/

Graduate bioinformatician/biochemistian with Statistic MA certification completing the final year of a Ph.D degree. Passionate about Big Data, Machine Learning and AI research, with strong technical and interpersonal skills, adept at working in teams and successfully delivering projects. I was awarded **Microsoft Azure Research Funding** in 2016 and gained a **Grow with Google Challenge Scholarship** in 2018.I am a certified **SAS** programmer and certified **Deep Learning** and **Cuda** instructor.

Education

| _ | University of Kentucky | Kentucky |
|---|--|---------------------------|
| 0 | Ph.D. in Biochemistry (Bioinformatics), | 2013–2019 |
| 0 | University of Kentucky Master (Cert) in Statistics | Kentucky 2016–2019 |
| 0 | Udacity Self-Driving Nano-degress (Expected) | 2018–2019 |

Technical Skills

- o Programming Languages: CUDA, C++ (OK), R, Python3, SAS and Matlab (OK). Please refer to:
 - Hackerrank: https://www.hackerrank.com/billchenxi
 - NVidia: https://nvidia.qwiklab.com/public_profiles/5521a192-c2e9-4899-9750-500959646159
- o AI/DL/RL/ML Knowledge: TensorFlow, PyTorch. Please refer to:
 - Courser: https://www.coursera.org/user/902cbf6a236e768db1c185c96778971b
- o Cloud Computing Skills: AWS, Google Cloud (GCP), Shiny. Please refer to:
 - **GCP/AWS**: https://qwiklabs.com/public_profiles/032b735c-3942-4f65-96e2-46bc821a884a

Publication & Software & Packages & Talks

- o Papers:
 - Automatic 13C Chemical Shift Reference Correction for Unassigned Protein NMR Spectra, Journal of Biomolecular NMR, Aug 2018 DOI: 10.1007/s10858-018-0202-5 (link)
 - Parallelized Interactive Machine Learning on Autonomous Vehicles, NAECON Dec 2018 DOI: 10.1109/NAE-CON.2018.8556776 (link)
 - Deep Learning by Doing: The Nvidia Deep Learning Institute, *Journal of Computational Science Education, Dec* 2018 DOI: 10.22369/issn.2153-4136/10/1/16 (link)
 - Pan-Cancer Epigenetic Biomarker Selection from Blood Sample Using SAS®, MWSUG, Sep 2018 (link)
 - Finite Mixture-of-Gamma Distributions: Estimation, Inference, and Model-Based Clustering, *Advances in Data Analysis and Classification, Jan 2019*
- o Package
 - BaMORC CRAN, 2018 CRAN GitHUB
 - BMRBr CRAN, 2017 CRAN GitHUB
- o Workshop
 - CUDA Programing Workshop (UK ACM 18)
 - Deep Learning for Computer Vision Workshop (UK ACM 19)

Notable Projects

- o (Deep Learning) Deep Learning for Cancer Classification with Gene Expression Data: Deep learning approach to classifying cancer types. Data from NIH Cancer Data public repo, with 32 different cancer types. The model accuracy is more than 97% with 0.2% false positive/negative rates.
- o (Reinforcement Learning) Parallelized Interactive Machine Learning on Autonomous Vehicles: Development of an interactive reinforcement learning pipeline for autonomous driving from a game simulation. The simulator is from Microsoft AirSim and driving environment is simulated from Unreal Engine.
- o (App Prototyping/Dev) CatsHub App Entrepreneurship: Development of student-oriented phone application that simplifies student's life by providing a one-stop shopping service hub. The original idea was from the University of Kentucky student organization. We pitched and created a prototype, and won venture reward.

Previous Employment

Deep Learning Institute

NVIDIA

University Ambassadors / Deep Learning Institute (DLI) Certified Instructor

2018-Present

Certified to teach Fundamentals of Computer Vision (CV), Fundamentals of Deep Learning for Multiple Data Types (MDT), and Fundamentals of Deep Learning for Natural Language Processing (NLP). Contents include: Cuda, Deep Learning, Neural Network, Image Classification, Object Detection, Image Segmentation, Word Generation, Image and Video Captioning, Text Classification, Text Translation, and etc..

Technologies: TensorFlow, Cuda, PyTorch, Caffe2, Keras, etc.

Moseley Bioinformatics Lab

Lexington KY

Graduate Research Assistant

2013-Present

Worked on construction of Protein NMR Reference Correction and Protein NMR Deuteration Level Detection frameworks. Designed and implemented the corresponding software package (BaMORC) and web applications systems. Created R package 'BMRBr' for easy download and processing of BMRB files.

Technologies: R, R Package, RStudio, Python, Shiny, Docker.

Prof. Derek S. Young, Dept. of Statistics, University of Kentucky Research Collaborator

Lexington KY

2017-Present

Worked on simulation for the Mix-Gamma Model with R. Jobs included generation of a large amount of simulated data from gamma distributions, implementation of the testing code, unit test, and preparation and design of submission shell code for HPC (High-performance Cluster).

Technologies: SAS, R, R Package, HPC Management, Bash

Reward and Certification Highlights

- o SIAM International Conference on Data Mining Dissertation Travel Award (2018)
- o MWSUG Conference Paper Scholarship (2018)
- o Grow with Google Challenge Scholarship (2017/18)
- o SC18 Paper Scholarship (2018)
- o SAS Certified Clinical Trials Programmer, Statistical Business Analyst, Advanced Programmer
- o Deeplearn.ai Certification Series (Coursera)