LONG WANG

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EDUCATION

Carnegie Mellon University (CMU) Pittsburgh, PA

May 2018

Master of Science in Computational Biology, GPA: 3.63

Huazhong Agricultural University (HZAU) Wuhan, China

Jul 2015

Bachelor of Science in Bioinformatics, Valedictorian, GPA: 3.68

WORK EXPERIENCE

SemanticMD, Inc. Pittsburgh, PA.

May 2017–Aug 2017

Deep Learning R&D Intern

• Took the lead on three deep learning pipelines development, including to detect sperm cells from cell cluster images and recognize their classes, to identify diabetic retinopathy patients, and to classify two-chambers vs. four-chambers ultrasound videos.

Jianbing Yan's Lab National Key Laboratory of Crop Genetic Improvement. Wuhan, China. Aug 2014–Jul 2015 Research Assistant (published one paper)

• Developed a tree-based machine learning method to detect SNP associations with rare variants in maize

Lingling Chen's Lab College of Informatics, HZAU. Wuhan, China.

Jul 2013-Jun 2016

Research Assistant (published three papers)

- Developed an ensemble model to infer transcriptional regulation using RNA-seq in Prokaryotes.
- Developed a statistical method to predict protein-protein interaction network and performed network analysis.

SELECTED COURSE PROJECTS

N-Gram probabilistic model (Java, HBase | Independent)

- Ran ETL on raw data and implemented a N-Gram language model using MapReduce to predict the next word
- Read plain text input file from HDFS, wrote statistical result to HBase, and connected the client interface with database to run phrases auto-completion.

Diabetic Retinopathy Classification with Imbalanced, Low-resolution retina images (Python | Independent)

- Designed and implemented a pipeline to do the classification (applied contrast limited adaptive histogram equalization (CLAHE) to enhance the contrast of the images, constructed a **GAN** to run data augmentation and applied transfer learning to avoid overfitting; **embedding resolution-aware** CNN with VGG16)
- Improved the accuracy from 92.9% (baseline) to 96.8%.

Machine learning in Large datasets (Python | Independent)

- Designed an efficient way to run a **scalable regularized logistic regression on** stochastic gradient descent (faster 10x than the classical LR model within a small memory)
- Implemented an **automatic reverse-mode differentiation** platform to train the CNN models (such as LSTM)
- Implemented a **memory-optimized** label propagation algorithm in Spark

3D Structure Reconstruction of Human Chromosomes from Hi-C data (Python | Team)

- Generated a **polynomial curve fit** between the interaction frequency (Hi-C data) and relative distance among nucleosome beads using the least squares method.
- Estimated the 3D coordinates for each nucleosome bead by posing the distance information to a constraint convex problem (find the minimal from the objective function using gradient descent)

Scrabble with Stuff (Java, JUnit Test | Independent)

- Designed a multiplayer Scrabble game from scratch using UML diagrams and design patterns
- Implemented core mechanics and the GUI of the game with Java Swing using MVC pattern
- Completed comprehensively unit tests using JUnit and Apache Ant

SKILLS

Programming: Java, Python, Scala, Bash, R, Matlab, Golang, Perl, C++.

Tools: Hadoop, Spark, Tensorflow, Pytorch, OpenCV, Hbase, MySQL, HDFS, MapReduce, Kafka, Scikit-learn.

ACADEMIC HONORS AND AWARDS

MSCB Merit Fellowship, Computational Biology Department. CMU, USA. (twice)	Aug 2017
Honorable Mention, Interdisciplinary Contest in Modeling, Comap Inc, USA.	Apr 2014
First Prize, Chinese Undergraduate Mathematical Contest in Modeling, Ministry of Education, CN.	Nov 2013
National Scholarship, Ministry of Education, China. (for top 0.5% students)	May 2012