# LONG WANG

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#### **EDUCATION**

Carnegie Mellon University (CMU) Pittsburgh, PA

May 2018(expected)

Master of Science in Computational Biology

Huazhong Agricultural University (HZAU) Wuhan, China

Jul 2015

Bachelor of Science in Bioinformatics. Valedictorian

### **WORK EXPERIENCE**

## SemanticMD, Inc. Houston, TX.

May 2017–Aug 2017

### Deep Learning R&D Intern

- Prepossessed on images and videos to filter out essential features using computer vision algorithms
- Developed and implemented machine learning pipelines to 1. detect and identify sperm cells; 2. identify diabetic retinopathy and glaucoma patients; 3. classify two-chambers vs. four-chambers ultrasound videos.

Jianbing's Laboratory National Key Laboratory of Crop Genetic Improvement. Wuhan, China. Aug 2014–Jul 2015 Research Assistant

Have published one paper as a participating author in *Nature Communication*.

**Lingling's Laboratory** *College of Informatics, HZAU.* Wuhan, China.

Jul 2013-Jun 2016

#### Research Assistant

Have published two papers as a participating author in *FEBS Letters* and *Scientific Reports*, respectively and a paper as the first author has been accepted in *Current Bioinformatics*.

#### **SELECTED COURSE PROJECTS**

# Diabetic Retinopathy Classification with Imbalanced, Low-resolution Retina Images

• Designed a pipeline to achieve high accuracy on diabetic retinopathy classification with imbalanced, low-resolution retina Images (contrast limited adaptive histogram equalization (CLAHE) to run image prepossessing; Resolution-aware CNN to enhance image resolution; GAN to run data augmentation; transfer learning to improve accuracy)

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

- Extracted interaction frequency information from Hi-C data
- Estimated the 3D position for each nucleosome bead by posing the distance information to a constraint convex problem (solved two objective functions via gradient descent)
- Visualized 3D structure results in PYMOL

# Machine learning in Large datasets (Independent)

- Applied sequential analysis on one-month Wikipedia data (set filters and MapReduce pipeline in AWS)
- Designed an efficient way to run regularized logistic regression on stochastic gradient descent in a cluster
- Implemented automatic reverse-mode differentiation for multilayer perceptron and long short term memory (LSTM) on large datasets in a cluster environment

# **Retrieval Algorithms for Search Engine** (Java | Independent)

- Implemented Boolean retrieval, BM25 and Indri ranking, and learning-to-rank capabilities.
- Implemented query expansion capabilities(pseudo-feedback) and diversification (PM-2 and xQuAD).
- Evaluate performance by calculating and comparing different metrics.

### **Twitter Analytics on the Cloud**

- Implemented Extract, Transform and Load (ETL) and streaming algorithm on more than 1 TB data
- Designed the databases structure to deal with scale to improve throughput.

### **SKILLS**

**Programming**: Java, Python, R, Perl, Matlab, Golang, C++, Shell.

Tools: Git, Hadoop, Spark, Scikit-learn, OpenCV, Tensorflow, Weka, GATK.

### **ACADEMIC HONORS AND AWARDS**

MSCB Merit Fellowship, Computational Biology Department. CMU, USA.

Honorable Mention, Interdisciplinary Contest in Modeling, Comap Inc. USA.

Aug 2017

Apr 2014

First Prize, Chinese Undergraduate Mathematical Contest in Modeling, Ministry of Education, China. Nov 2013

National Scholarship, Ministry of Education, China. (for top 0.5% students)

May 2012