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#### I. INTERESTS

Big Data, Cloud Computing, Biological Computing, Machine Learning, Deep Learning

#### II. QUALIFICATION

- In-depth understanding and rich hands-on experience in Hadoop, Spark, Hive, Pig, and Tez frameworks.
- Strong parallel programming experience in MPI, Spark, Hadoop MapReduce, Pig Latin Data Flow, and HiveQL Language.
- Proficient in shell, SQL programming
- Hands-on experience in big data analysis, large scale biological data processing.
- 6 years working experience as a software engineer in multinational project teams.

# III. EDUCATION

• 08/2015 ~ present	PhD Candidate	Computer Science	GPA 4.0	Florida State University, FL
• $08/2014 \sim 08/2015$	M.S.	Computer Science	GPA 4.0	Auburn University, AL
<ul> <li>08/2004 ~ 06/2007</li> </ul>	B.S.	Computer Science	North China	Electric Power University, China

#### IV. EXPERIENCE

•	Teaching Assistant	08/2015 ~ present	Florida State University		
•	Summer Intern	$05/2015 \sim 08/2015$	Lawrence Berkeley National Laboratory		
•	Research Assistant	$08/2014 \sim 05/2015$	Auburn University		
	Thesis: Feature Enhancement and Performance Evaluation of BioPig Analytics				

# V. PROJECTS

# At Florida State University:

- Designed a hybrid clustering algorithm based on LPA for metagenome read clustering
- Developed a scalable sequence clustering tool named SpaRC based on Apache Spark.
- Designed a cross-layer scheduler for improving DAG-structured query processing in MapReduce.

### At Lawrence Berkeley National Lab:

- Upgraded BioPig toolkit from Hadoop 1 to Hadoop 2 (2.7.0).
- Tuned Hadoop parameters for BioPig performance optimization.
- Implemented K-mer Similarity to extended BioPig toolkit functionality

### At Auburn University:

- Implemented an MPI program to sort millions of integers using multiple algorithms.
- Conducted research on two big data frameworks: Hadoop and Spark.
- Developed an android project that can support downloading files from top cloud storage providers.

# At PwC, CleNET and Kingbase:

• Developed/Maintained projects in various domains using C#, Java languages and SQL server databases.

# VI. PUBLICATIONS

- Lizhen Shi, Zhong Wang, Weikuan Yu, Xiandong Meng. *Performance Evaluation and Tuning of BioPig for Genomic Analysis*. International Workshop on Data-Intensive Scalable Computing Systems (DISCS) in conjunction with the ACM/IEEE Supercomputing Conference (SC'15), Austin, TX, November 2015.
- Lizhen Shi, Zhong Wang, Weikuan Yu, Xiandong Meng. A Case Study of Tuning MapReduce for Efficient Bioinformatics in the Cloud. Journal of Parallel Computing (ParCo'16).
- Ji Huang, Stefania Vendramin, **Lizhen Shi**, Karen M McGinnis. *Construction and optimization of a large gene Co-expression network in maize using RNA-Seq data*. Plant physiology (2017)
- Lizhen Shi, Xiandong Meng, Elizabeth Tseng, Michael Mascagni, Zhong Wang. SpaRC: scalable sequence clustering using Apache Spark. Bioinformatics.

- Lizhen Shi, Volkan Sevim, Michael Mascagni, and Zhong Wang. Leveraging long-read sequencing for cost-effective metagenome clustering. (To be submitted)
- Lizhen Shi, Bo Chen. A Vector Representation of DNA Sequences Using Locality Sensitive Hashing. (bioRxiv)