

Zhen Zhang

Curriculum Vitae

Education

2010–2016 **Doctor of Philosophy in Bioinformatics**, *Central South University*, Changsha, China *Grade*

My research interests include structural variation detection, read mapping and genome assembly.

2001–2006 **Master of Science in Computer Science**, *Technical University Munich*, Munich, Germany, *GPA – 2.3* (1.0 is the best grade).

I specialized in service-oriented architecture and full stack web development.

1996–2000 **Bachelor of Engineerer in Environmental Engineering and Technology**, *Hunan University*, Changsha, China, *GPA – 3.9*.

I earned a minor in Computer Science.

Experience

Vocational

2009–present Lecturer, Hunan Institute of Science and Technology, Yueyang, China.

I teached the following two courses: Software Engineering and Algorithm Design and Analysis.

2007–2008 **Junior Developer**, *Computershare Deutschland GmbH & Co. KG*, Munich, Germany.

I was in charge of three projects: 2publish customisation, Hewlett Packard Promotool and eCampaigns.

2005–2007 **Programmer Intern**, *Intergraph Deutschland GmbH*, Ismaning, Germany.

I automated the setup package building and deployment process for a GIS web application flimasWeb. Besides, I conducted a performance test for flimasWeb.

Miscellaneous

2004–2005 **Student Trainee**, *Rechts der Isar Hospital*, Munich, Germany.

I developed a serial C++ driver for the Aurora magnetic tracker under Linux.

Award

2015 Best Paper Award, Joint Genome Informatics Workshop (GIW) and International Conference on Bioinformatics (InCoB), Tokyo, Japan, Sept. 9-11, 2015

Languages

Chinese Native speaker

English **Professional working proficiency**German **Professional working proficiency**

Computer skills

Intermediate Git, LATEX, Microsoft Visio, Javascript

Advanced C++, Python, Java, Full Stack Web Developement, Linux

Interests

- Mountain climbing

- Traveling

- Stamp collecting

- Coin collecting

Publications

[1] Zhen Zhang, Jianxin Wang, Junwei Luo, Xiaojun Ding, Jiancheng Zhong, Jun Wang, Fang-Xiang Wu, and Yi Pan. Sprites: detection of deletions from sequencing data by re-aligning split reads. *Bioinformatics*, Feb 2016. doi: 10.1093/bioinformatics/btw053. URL http://dx.doi.org/10.1093/bioinformatics/btw053.