#### (Info)

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## **Education**

2010.08 - 2015.07 PhD. Academy of Military Medical Sciences. *Bioinformatics* 

2006.08 - 2010.07 BSc. Tsinghua University. Computer Science and Technology

# **Experience**

2015.08 - Present Assistant Researcher. Academy of Military Medical Sciences

## [Awards]

2011 - 2014 Postgraduate Scholarships in Academy of Military Medical Sciences

2015 Excellent Doctor Degree Dissertation in Academy of Military Medical

Sciences

#### [Research Area]

Integrated analysis of multi-omics, big data and biological network models.

#### **Skills**

- 1. Excellent understanding of bioinformatics, human genomics and different genome sequencing platforms. More than 5 years' experience with omics data generated from next generation sequencing. Familiar with common databases and tools.
- Good knowledge of C++, Jave, scripting languages (Perl, Python) and mathematics tools (R, Matlab). Basic knowledge of MySQL. Experience in system management in a Linux environment.

- 3. Fluent in reading, writing and speaking Chinese and English.
- 4. Excellent communication and interpersonal skills, team working and self-driven.

### [Publications]

- 1. **Shen W**<sup>1</sup>, Wang D, Ye B, Shi M, Ma L, Zhang Y\*, Zhao Z\*. GC3-biased gene domains in mammalian genomes. *Bioinformatics*. 2015, 31(19):3081-3084. (IF=5.8)
- 2. **Shen W**<sup>1</sup>, Wang D, Ye B, Shi M, Zhang Y\*, Zhao Z\*. A possible role of Drosophila CTCF in mitotic bookmarking and maintaining chromatin domains during the cell cycle. *Biological Res.* 2015, 48:27. (IF=1.3)
- 3. Zhang Y<sup>1</sup>, **Shen W**<sup>1</sup>, Shi M<sup>1</sup>, Zhang L, Zhang Z, Li P, Xing LY Luo F, Sun Q\*, Zheng X, Yang X\*, Zhao Z\*. Involvement of aberrant miR-139/Jun feedback loop in human gastric cancer. *Biochim Biophys Acta*. 2015, 1853(2):481-488. (Co-first author, IF=5.3)
- 4. Ye B<sup>1</sup>, **Shen W**, Wang D, Li P, Zhang Z, Shi M, Zhang Y, Zhang F\*, Zhao Z\*. ZNF143 is involved in CTCF-mediated chromatin interactions by cooperation with cohesin and other partners. *Mol Biol*. 2016, 50(3):496-503. (IF=0.6)
- 5. Zhang Z<sup>1</sup>, Zhang Y, Shi M, Ye B, **Shen W**, Li P, Xing L, Zhang X, Hou L, Xu J\*, Zhao Z\*, Chen W\*. Anthrax Susceptibility: Human Genetic Polymorphisms Modulating ANTXR2 Expression. *Toxins*. 2016, 8(1). (IF=3.6)
- 6. Zhang Y<sup>1</sup>, Fan K<sup>1</sup>, Sun Q<sup>1,\*</sup>, Chen A, **Shen W**, Zhao Z, Zheng X<sup>\*</sup>, Yang X<sup>\*</sup>. Functional screening for miRNAs targeting Smad4 identified miR-199a as a negative regulator of TGF-β signalling pathway. *Nucleic Acids Res.* 2012, 40(18):9286-9297. (IF=9.2)