# Ting He

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### **Education**

#### **Bachelor of Science in Bioinformatics**

Dalian University of Technology(project 211 and 985) GPA: 3.5/4.0 (top 6) Major GPA:3.8/4.0 (top 1)

#### **Main Courses:**

- Data Structures and Algorithms
- Database Principles
- Bio Java

- Systems Biology
- Genomics
- Molecular biology
- Biostatistics
- Data Mining in Bioinformatics
- Perl Programming
- Computational Biology
- Proteomics
- Cell Biology
- Biochemistry

### **Publications**

Zhenyao Zhang, **Ting He**, *Comparative proteomics analysis of human osteosarcoma by 2D DIGE with MALDI-TOF/TOF MS*, The First Youth Medical Forum of Chinese Society of Biochemistry and Molecular Biology, March 2016

## Research experience

### Visiting Student, University of Calgary, Canada

September-December 2016

Frequent Itemset Mining, in collaboration with Dr. Quan Long

- Discretized continuous data based on statistic model
- Used machine learning methods to advance the investigation of phenotype-genotype relationships in HDD

## Visting Student, University of Calgary, Canada

August-September 2016

Cancer Genome data analysis, in collaboration with Dr. Quan Long and Lauren Mak

- Used softwares to call SNPs in next-generation sequencing data
- Compared quality and accuracy of SNP calling in different pipelines

### Participator, Dalian University of Technology, China

**April-September 2015** 

Research of synthesis of common drugs

• In charge of improving synthesis route for ciprofloxacin

#### **Project Experience**

# Team lead, Designer and Programmer, Dalian University of Technology, China

**July-September 2015** 

Health Data Analysis app development

- Led a team of 6 students to develop an app for patients to enter health data, and app would make general recommendations to improve health
- Designed an attractive user interface and functionality developments

#### **Executor, Dalian University of Technology, China**

**July-August 2015** 

Drug molecular design

- Investigated interaction of ligands and protein using molecular simulation method
- Created and profiled pharmacophore and then used virtual screening

### **Work Experience**

#### August 2016 – Present, Bioinformatics Skills internet forum

Manager, Writer, Editor, website: http://www.biotrainee.com/

- Worked with various top schools' masters, phds and bioinformatics hobbyists
- Educating biologists in bioinformatics skills and pipelines

### Honors and awards

- First-class, China National Scholarship (0.2% among all university students in China), Ministry of Education of the P. R.C, October 2016
- First prize(Top 1/50)/second prize(Top 10/50), Scholarships for Excellent Performance, Dalian University of Technology, 2015 and 2016
- First Prize(Top10/1000), Scholarships for Technological Innovation, Dalian University of Technology, 2015
- First prize, The 12th Climbing Competition(for technological achievement), Dalian University of Technology, 2016
- Third prize, Mobile Application Development, Liaoning Provincial Education Authourity, 2015
- Second prize, Mathematical modeling competition of China Certification cup, Global Mathematical Modeling Certificate Authority, 2016
- The honor of All-around Good Student(for academic achievement and community involvement), Dalian University of Technology, 2015 and 2016

## **Skills**

	Programming languages		Software experience		Laboratory experience	
•	Java	•	BWA	•	Cell culture	
•	Perl	•	Samtool	•	Inverse PCR	
•	C++	•	LibDock	•	Agar-gel electrophoresis	
•	MySQL	•	AMBER software tools	•	Heat shock method	
•	Matlab	•	PBS/TQRUE			