Shoujun Gu, Ph.D

Rockville, Maryland

Email: *shoujun.gu@gmail.com* | Tel: 210-315-2598

Website: *https://shoujungu.github.io/introduction/*

https://www.linkedin.com/in/shoujun-gu-bb624643/

Technical Skills

• *Programing Language:*
Python: Numpy, Pandas, Matplotlib, Seaborn, SQLAlchemy, Flask, Scikit-learn, Keras
R: DESeq, EdgeR, maSigPro

• *Interactive Data Visualization & Webpage:*
Javascript: D3js, Plotly, Leaflet
HTML: Bootstrap

• *Database:*
MySQL, MongoDB

• *Bioinformatics & Next Generation Sequencing:*
Data mining from various biology databases: NCBI, Ensembl, TCGA, cBioportal
Hands-on experience in NGS data analysis workflow

• *Molecular & Cellular Biology:*
Extensive experience in most molecular & cellular experiment, as well as animal model

Data Analytics Projects

• *The Cancer Genome Atlas (TCGA) data mining and analysis project*
The manuscript of this project entitled '*A Pan-Cancer Atlas of Genomic, Epigenomic and Transcriptomic Alterations in the TGF-β Pathway*', is under reviewing.

• *Built a machine learning model to predict the significance of a research project based on its abstract*
Details available at: *https://github.com/shoujungu/Impact_Factor_Pred*
Sample app available at: *https://if-pred.herokuapp.com*

• *Interactive visualization of US healthcare providers data*
Details available at: *https://github.com/shoujungu/US_Healthcare_Providers*
Sample app available at: *http://hcproviders.herokuapp.com*

• *Data mining and analysis on all healthcare publications (Year: 2017) in PubMed database*
Details available at: *https://github.com/shoujungu/2017_Pubmed_Analysis*

Working Experience

The George Washington University, Washington, DC Oct 2015 – Jan 2018
Postdoctoral Researcher

- Discovered an aberrant regulation pathway between TGF-β signaling and cancer biomarker CEA in colon adenomas by using both whole genomic sequencing and whole transcriptome sequencing methods [1];
- Discovered that Vitamin D promotes liver tumor progression in TGF-beta deficient environment by using both in vivo mouse model and bioinformatics tools [2];
- Published a review article entitled "Alcohol, stem cells and cancer" [3];
- TCGA data mining and analyzing of Pan-Cancer Atlas in the TGF-β signaling (under review).

National Institutes of Health, Bethesda, MD Oct 2012 – Sept 2015
Visiting Fellow

- Discovered that Wnt5a-induced Vangl2 phosphorylation is a key step for planar cell polarity (PCP) signaling initiation and establishment by showing phosphorylated Vangl2 exhibits altered interaction with other PCP proteins and forms self-aggregation;
- Generated myc-Prickle1 knock-in mice by using CRISPR/Cas9 genome editing technique;
- Discovered that Casein Kinase 1 ε/δ conditional knockout mice exhibit significantly shorted long bones and delayed limb development.

Education

The George Washington University, Washington, DC Oct 2017 – Apr 2018
Data Analytics Boot Camp

An intensive 24-week long boot camp dedicated to Data Mining and Analytics Skills on a variety of real-world problems.

UT Health Science Center at San Antonio Aug 2005 – May 2012
Ph.D. Department of Biochemistry

- Oral presentation in American Society for Bone and Mineral Research 2008 Annual Meeting;
- Discovered that the Twist1 inhibits Sox9 transactivation through protein-protein interaction between Twist1 C-terminal domain and Sox9 DNA binding domain, which results in negative regulation of chondrogenesis [4].

Publications

1. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4839765/
2. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4960540/
3. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC5724803/
4. https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3375531/