Xiaojie Qiu, PhD Candidate

CONTACT Information Department of Genome Sciences

MCB program

University of Washington

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 $O \\ \text{BJECTIVE}$

To obtain a postdoctoral position at the interface of computational biology and single-cell genomics.

РнD

University of Washington

Aug 2013 – Jun 2018 (Expected)

Molecular and Cellular Biology (MCB) program

- Reconstructing developmental trajectories with single-cell RNA-seq
- Inferring causal regulatory networks underlying cellular fate transitions

Masters

East China Normal University, BIOINFORMATICS

Aug 2009 - Jun 2012

• Mathematical modeling of cell differentiation, transdifferentiation and reprogramming

BA

Changchun University of Technology, BIOENGINEERING

Aug 2004 – Jun 2008

RESEARCH EXPERIENCE University of Washington, Seattle, WA

tle, WA Aug 2013 – Present

Graduate Research Assistant (Advisors: Dr. Cole Trapnell)

Institute for Systems Biology, Seattle, WA Research Staff (Advisors: Dr. Sui Huang)

Shanghai Jiaotong University, Shanghai, China

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Sep. 2011 – Jun. 2012

Jan. 2012 - Sep. 2013

Research Assistant (Advisors: Dr. Ping Ao)

East China Normal University, Shanghai, China Master Candidate Research Assistant (Advisors: Dr. Tieliu Shi) Sep. 2009 – Jun. 2012

Honors and Awards China Scholarship Council Award (Finalist)

2017

This award honors overseas Chinese students with outstanding academic accomplishments from different disciplines.

JOURNAL PUBLICATIONS

- X Qiu, Q Mao, Y Tang, L Wang, R Chawla, H Pliner, C Trapnell. "Reversed graph embedding resolves complex single-cell trajectories." Nature methods, ADVANCE ONLINE PUBLICATION, 2017. doi:10.1038/nmeth.4402
- [2] X Qiu, A Hill, J Packer, D Lin, YA Ma, C Trapnell "Single-cell mRNA quantification and differential analysis with Census.", Nature methods 14 (3), 309-315, 2017. doi:10.1038/nmeth.4150
- [3] J Cao, J S. Packer, V Ramani, D A. Cusanovich, C Huynh, R Daza, X Qiu, C Lee1, S N. Furlan, F J. Steemers, A Adey, R H. Waterston, C Trapnell, J Shendure. "Comprehensive single-cell transcriptional profiling of a multicellular organism." Science 357.6352: 661-667, 2017. DOI: 10.1126/science.aam8940.
- [4] NK Hanchate, K Kondoh, Z Lu, D Kuang, X Ye, X Qiu, L Pachter, C Trapnell, L B Buck. "Single-cell transcriptomics reveals receptor transformations during olfactory neurogenesis." Science 350 (6265), 1251-1255, 2015. DOI:10.1126/science.aad2456
- [5] **X Qiu**, S Ding, T Shi. "From understanding the development landscape of the canonical fate-switch pair to constructing a dynamic landscape for two-step neural differentiation." **PloS one** 7 (12), e49271, **2012**. doi:10.1371/pone.0049271
- [6] J Wang, X Qiu, Y Li, Y Deng, T Shi. "A transcriptional dynamic network during Arabidopsis thaliana pollen development". BMC systems biology 5 (3), S8, 2011. doi.org/10.1186/ 1752-0509-5-S3-S8
- [7] B He, X Qiu, P Li, L Wang, Q Lv, T Shi. HCCNet: an integrated network database of hepatocellular carcinoma. Cell research 20 (6), 732, 2010. DOI:10.1038/cr.2010.67

Submitted

- [8] D Cacchiarelli, X Qiu, S Srivatsan, M Ziller, E Overbey, J Grimsby, P Pokharel, K Livak, S Li, A Meissner, T Mikkelsen, J Rinn, C Trapnell "Aligning single-cell developmental and reprogramming trajectories identifies molecular determinants of reprogramming outcome", 2017 (submitted). https://www.biorxiv.org/content/early/2017/03/30/122531
- [9] D A Cusanovich, JP Reddington, DA Garfield, R Daza, R Marco-Ferreres, L Christiansen, X Qiu, F Steemers, C Trapnell, J Shendure, EEM Furlong. "The cis-regulatory dynamics of embryonic development at single cell resolution", 2017 (Under review in Nature). https://www.biorxiv. org/content/early/2017/07/20/166066
- [10] H Pliner, J Packer, J McFaline-Figueroa, D Cusanovich, R Daza, S Srivatsan, X Qiu, D Jackson, A Minkina, A Adey, F Steemers, J Shendure, C Trapnell. "Chromatin accessibility dynamics of myogenesis at single cell resolution", 2017 (Under review in Cell). https://www.biorxiv.org/content/early/2017/06/26/155473

BOOK CHAPTERS

[11] JX Zhou, X Qiu, AF d'Hroul, S Huang. "Discrete Gene Network Models for Understanding Multicellularity and Cell Reprogramming: From Network Structure to Attractor Landscape." in Computational Systems Biology, 2rd ed., Elsevier Inc. 2013.

SOFTWARE PRODUCTS

- [12] C Trapnell, D Cacchiarelli, X Qiu. "Monocle". http://cole-trapnell-lab.github.io/monocle-release/. 2017.
- [13] T Lin, S Hughes, X Qiu. "densityClust". https://github.com/Xiaojieqiu/densityClust. 2017.
- [14] X Qiu, C Trapnell, Q Mao, L Wang. "DDRTree". https://cran.r-project.org/web/packages/DDRTree/index.html. 2017.
- [15] X Qiu, C Trapnell, Q Mao, L Wang. "SimplePPT". https://github.com/cole-trapnell-lab/monocle2-rge-paper/tree/master/Supplementary_scripts/Packages. 2017.
- [16] X Qiu, C Trapnell, Q Mao, L Wang. "L1Graph". https://github.com/cole-trapnell-lab/monocle2-rge-paper/tree/master/Supplementary_scripts/Packages. 2017.

SCIENTIFIC COMPUTING SKILLS

Languages Proficient in R, bash/csh, Python, Julia; Familiar with C++, PHP, HTML and others.

Other Tools Proficient in MatLab, Cytoscape, Illustrator, git; Familiar with Languages Proficient in MatLab, Cytoscape, Illustrator, git; Familiar with Languages Proficient in MatLab, Cytoscape, Illustrator, git; Familiar with Languages Proficient in R, bash/csh, Python, Julia; Familiar with C++, PHP, HTML and others.

EXPERIMENTAL SKILLS

Tissue cell culture, RNA-seq library preparation, Flow cytometry, other basic molecular biology techniques

Outreach and Teaching

Science Education Partnership, FRED HUTCHISON CANCER RESEARCH CENTER Fall 2016 Tutor. Trained middle school teachers in molecular biology techniques and how to apply them in laboratory research.

STEMPREP Program, University of Washington

Summer 2017

Tutor. The STEMP prep program aims to assist in producing the next generation of minority researchers in Science, Technology, Engineering, Math and Medicine (STEMM).

References

PhD Thesis Advisor: Cole Trapnell, PhD

Assistant Professor Genome Sciences Department http://cole-trapnell-lab.github.io/ coletrap@uw.edu Jay Shendure, MD, PhD Professor, HHMI investigator Genome Sciences Department http://krishna.gs.washington.edu/ shendure@uw.edu

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