# Sarah A. Christensen

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## EDUCATION

UNIVERSITY OF ILLINOIS

PhD Candidate in **Computer Science** Advised by Mohammed El-Kebir and Tandy Warnow

- My research is in the intersection of theoretical computer science and computational biology; in
- particular, I develop efficient algorithms to support biological and oncological research.
- Cumulative GPA: 3.95/4.0; Chirag Fellowship (2016-2018), State Farm Doctoral Award (Spring 2018), C.L. and Jane Liu Award (Spring 2018), Ira & Debra Cohen Graduate Fellowship (Spring 2019).

#### AMHERST COLLEGE

Bachelor of Arts with Distinction

- Major: Mathematics; Cumulative GPA: 3.83/4.0; Major GPA: 3.97/4.0
- Member of The Phi Beta Kappa Society; Top 10% of graduating class

# WORK & RESEARCH EXPERIENCE

#### MCKINSEY & COMPANY

Summer Associate

• Collaborated with clients and vendors to build climate change modelling capabilities that predict various asset risks.

#### KALLYOPE

Computational Genetics Research Intern

• Designed a novel graph algorithm to efficiently delineate correlated regions of the human genome within finite populations. Implemented a parallelized version of the algorithm in C++. Accepted talk at ASHG with journal publication forthcoming.

#### **CORNERSTONE RESEARCH**

Economic Consulting Senior Analyst

• Structured open-ended analytical tasks, synthesizing the results into intuitive visual exhibits. Presented results at client and expert meetings. Managed junior analysts in data collection and econometric analysis.

#### MASSACHUSETTS GENERAL HOSPITAL

Research Assistant in the Division of General Medicine

• Analyzed the cost-effectiveness of deploying a new assay technology in antenatal clinics using Monte Carlo simulations.

### YALE SCHOOL OF PUBLIC HEALTH

Research Assistant to Professor Alison Galvani

• Performed meta-analysis on key influenza transmission model parameters to access robustness of modeling predictions.

# PUBLICATIONS

- Christensen S., Kim J., Chia N., Koyejo O., El-Kebir M. (2020) Detecting Evolutionary Patterns of Cancers using Consensus Trees. *Proceedings of the European Conference on Computational Biology (ECCB)*.
- Yu X., Le T., Christensen S., Molloy E.K., Warnow T. (2020). Advancing Divide-and-Conquer Phylogeny Estimation using Robinson-Foulds Supertrees. *Proceedings of the 20th International Workshop on Algorithms and Bioinformatics*.
- Christensen S., Leiserson M.D.M., El-Kebir M. (2020). PhySigs: Phylogenetic Inference of Mutational Signature Dynamics. *Proceedings of the Pacific Symposium on Biocomputing (PSB)*.
- Christensen S., and McManus J. (2019) Universal LD Blocks in the Human Genome. Accepted platform talk at the 69th Annual Meeting of *The American Society of Human Genetics (ASHG)*. Houston, Texas.
- Christensen S., Molloy E.K., Vachaspati P., Warnow T. (2019) TRACTION: Fast Non-Parametric Improvement of Estimated Gene Trees. *Proceedings of the 19th International Workshop on Algorithms and Bioinformatics*.
- Christensen S., Molloy E.K., Vachaspati P., Warnow T. (2018) OCTAL: Optimal Completion of Gene Trees in Polynomial Time. *Algorithms for Molecular Biology*.
- Christensen S., Molloy E.K., Vachaspati P., Warnow T. (2017) Optimal Completion of Incomplete Gene Trees in
  Polynomial Time using OCTAL. *Proceedings of the 17th International Workshop on Algorithms and Bioinformatics*.
- Ciaranello AL, Myer L, Kelly K, Christensen S, Daskilewicz K, et al. (2015) Point-of-Care CD4 Testing to Inform Selection of Antiretroviral Medications in South African Antenatal Clinics: A Cost-Effectiveness Analysis. *PLoS ONE*.

Urbana-Champaign, Illinois

Amherst, Massachusetts September 2009 – May 2013

August 2016 - Present

New York, New York May 2020–July 2020

New York, New York August 2018–December 2018

New York, New York

July 2013 - July 2016

**Boston, Massachusetts** May 2012–August 2012

New Haven, Connecticut

May 2011–August 2011