# Danielle "Dasha" Pruss

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## Education

BS, Computer Science, *University of Utah* Dean's List, *2011-2015*  2016

# Research Experience

## Research Assistant, University of Utah

2015 - present

Working under Dr. Miriah Meyer in the Vis Design Lab of the Scientific Computing and Imaging Institute. Visualizing the retinal cell connectome, in collaboration with the Marc Laboratory of the Moran Eye Center. Research funding from the Scientific Computing and Imaging Institute.

## Research Intern, Harvard-MIT Health Sciences and Technology

2014

Worked under Dr. Alexander Gimelbrant of the Dana Farber Cancer Institute, an affiliate of Harvard Medical School. Performed bioinformatic analysis of gene expression variation in genes subject to monoallelic expression. Research funding from Harvard-MIT Health Sciences and Technology Summer Institute in Bioinformatics and Integrative Genomics.

## Research Technician, University of Utah

2012 - 2013

Worked under Dr. Michael Kay of the Department of Biochemistry. Synthesized and purified peptides for Ebola virus inhibition. Developed bioinformatics Perl scripts for mutant analysis of deep sequencing data. Research funding from Undergraduate Research Opportunities Program and the Biochemistry Department of the University of Utah.

# Internships

#### Software Engineering Intern, Qualcomm

2015

Worked on the QaFAST team (Qualcomm Application Framework for Automated Suite of Tests). Developed Android OS test automation frameworks for product level functional, performance, and stability testing. Revamped the Android test automation application user interface.

## Clinical Variant Research Intern, Myriad Genetics

2010 - 2011

Performed genetic and pedigree analysis of deleterious mutations and variants of uncertain significance in tumor suppressor genes BRCA 1/2, associated with increased risk of breast and ovarian cancer.

# **Teaching Assisting**

Leading weekly lab sessions, holding office hours, and grading.	
CS 2420, "Data Structures and Algorithms" Miriah Meyer, 1 semester	2015
CS 1410, "Object Oriented Programming," Embedded Systems Version Peter Jensen, 1 semester	2014
CS 1410, "Object Oriented Programming" Erin Parker, 2 semesters	2013 – 2014
Publications	
"Risk alleles in genes with monoallelic expression are enriched among gain- of-function variants and depleted among loss-of-function variants in neuro- developmental disorders." Savova V., Pruss D., Gimelbrant A., Weiss L.	in revision
"Statistical Soft Mutagenesis, a Tool for Peptide Phage Display Affinity Maturation as Applied to D-peptide Inhibitors of Ebolavirus Entry" Clinton T., Szabo-Fresnais N., Apple S., Pruss D., Pandya M., Whitby F., McKinnon R., Hill C., Welch B., Eckert D., Kay M.	in progress
Presentations	
"Expression Variation in Monoallelic Genes"	2014
2014 Harvard-MIT Health Sciences and Technology Summer Scholars Bioinformatics and Integrative Genomics, Harvard Medical School	Program in
Awards	
scholarships	
Hyde Merrill Endowed Scholarship  University of Utah, College of Engineering	2015
School of Computing Women's Scholarship University of Utah, School of Computing	2015
Grace Murray Hopper Memorial Scholarship University of Utah, School of Computing	2015
Honors at Entrance Scholarship  University of Utah, Honors College	2011 – 2015
university honors	
Dean's List University of Utah	2011 – 2015

## conference grants

2015 Grace Hopper Scholar (declined) Funding from the Anita Borg Institute	2015
2014 Grace Hopper Celebration of Women in Computing Scholarship Funding from the University of Utah, School of Computing	2014
Qualcomm Women's Collegiate Conference Scholarship Funding from the University of Utah, School of Computing	2014
2013 Rocky Mountain Conference for Undergraduate Women in Physics Funding from the University of Utah, College of Science	2013
research funding	
University of Utah, Scientific Computing and Imaging Institute	2015
Harvard-MIT Health Sciences and Technology, Summer Institute in Bioinformatics and Integrative Genomics	2014
University of Utah, Biochemistry Department	2012 – 2013
Undergraduate Research Opportunities Program (UROP)	2012
other	
Stanford University Summer Institute in Microbiology and Biotechnology	2010
Relevant Coursework and Projects	

## computer science

Perception for Graphics (CS 5650)

Data Visualization (CS 5630)

Interactive PAC12 Football Games visualization – Javascript, D3

Interactive United Nations MY World 2015 vote project visualization – Javascript, D3 Interactive Needle Electroanatomical Potential Tracking visualization (term project)

Database Systems (CS 5530)

Online bookstore: user statistics, book searches, purchases, and user recommendations – JSP and MySQL

Machine Learning (CS 5350)

Genetic variants for identifying cancer type (term project)

Implemented a Decision Tree through ID3 algorithm and 6-fold cross-validation – Java

Implemented Perceptron algorithm and batch/margin variants – Java

Implemented Support Vector Machine – Java

Natural Language Processing (CS 5340)

Implemented a probabilistic n-grams model of sentences – Java

Designed and built a question answering system using CBC's current event stories – Python

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Computer Security (CS 4964)

Computer Systems (CS 4400)
Implemented a buffer overflow attack – C
Developed a simple Unix shell – C
Implemented the functions malloc/realloc/free using segregated free lists – C

Algorithms (CS 4150)

Computer Organization (CS 3810)

Software Practice I/II (CS 3500, CS 3505)
Boggle game + user interface with score database – C# and MySQL
Multi-user spreadsheet with concurrent editing using socket implementations – C# and C++
FFmpeg image codec development, including encoding, compression and decoding – C

Models of Computation (CS 3100)

Data Structures and Algorithms (CS 2420)
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## life sciences

Genetics (BIOL 2030)

General Chemistry I/II (CHEM 1210/1220)

Organic Chemistry I/II (CHEM 2310/2320)

## math

Engineering Probability and Statistics (CS 3130)

Calculus I/II/III (MATH 1210/MATH 1220/MATH 2210)

Discrete Structures (CS 2100)

## writing

Professional Writing (WRTG 3015)

Scientific Writing (Summer Institute in Bioinformatics and Integrative Genomics)

## **Programming Languages and Environments**

# Languages

object oriented
Java, Python, C#, C++
web/data visualization
JavaScript, D3, HTML/CSS
database systems
MySQL
multi-paradigm
Perl, R, Racket
hardware
C, x86, Arduino (C/C++)

#### **Environments**

- Eclipse
- Visual Studio 2012/2013
- DrRacket
- RStudio
- JetBrains WebStorm, PyCharm, IntelliJ
- MySQL Workbench
- Linux environments (emacs, vi)

## **Version Control**

- Git
- Mercurial