# Arun Durvasula

@arundurvasula

Department of Plant Sciences Phone: (408) 656-6358

University of California Email: adurvasula@ucdavis.edu

1 Shields Ave. Blog: www.arundurvasula.wordpress.com Davis, CA 95616 Github: www.github.com/arundurvasula

#### Education

BS Biotechnology, Microbiology and Fermentation, University of California Davis 2015 (expected)

High School Diploma, Saint Francis High School, Mountain View, CA 2011

## Experience/Employment

Research Intern, Ross-Ibarra Lab. University of California, Davis. June 2013 - Present

Research Intern, Hancock Lab. Max F. Perutz Labratories (planned). June 2015 - August 2015

Research Assistant, Rowhani Lab. University of California, Davis.

June 2014 - June 2015

Technical Reviewer for Bioinformatics Data Skills, O'Reilly Media. December 2013 - March 2014

Principal Programmer, Seqcoverage. July 2014

Research Assistant, Tagkopulous Lab. University of California, Davis. January 2013 - June 2013

## Teaching

Teaching assistant: Ecological Genomics (Graduate), Winter 2015

#### Presentations and Posters

Patterns of Demography and Selection Since Maize Domestication, Poster at Maize Genetics Conf., 2015

ANGSD-wrapper: scripts to streamline and visualize NGS population genetics analysis, Poster at Bay Area Population Genetics Conf, 2014 and Maize Genetics Conf, 2015

#### **Publications**

Tyler Kent, Siddartha Bhadra-Lobo, **Arun Durvasula**, Jinliang Lang, Eric Fuchs, Jeffrey Ross-Ibarra. Population genomic assessment of crop-wild gene flow in the endangered wild rice *Oryza glumaepatula* (2015). In preparation.

**Arun Durvasula**, Tyler Kent, Jeffrey Ross-Ibarra. ANGSD-wrapper: scripts to streamline and visualize NGS population genetics analysis (2015). In preparation.

Timothy Beissinger, Li Wang, Matthew Hufford, **Arun Durvasula**, Jeffrey Ross-Ibarra. Patterns of Demography and Selection Since Maize Domestication (2015). In preparation.

Arun Durvasula 2

### Skills

Programming: R, Python, Bash, Awk, C, C++, Ruby, Java, Javascript, HTML, CSS

Frameworks/Libraries: Scikit Learn, Scipy, Numpy, Matplotlib,

Bioinformatics: assembly, alignment, population genetics analysis

Tools: Git, Shiny, Matlab, SQL, Machine learning, IPython, LaTeX, Slurm, Sun Grid Engine

Molecular biology: PCR, Gel electrophoresis, RFLP, Bacterial transformation, Genomic DNA isolation

### References

#### Jeffrey Ross-Ibarra

Associate Professor

Dept. of Plant Science

University of California

Davis, CA. 95616

email: rossibarra@ucdavis.edu

#### Maher Al Rwahnih

Project Scientist

Foundation Plant Services

University of California

Davis, CA. 95616

email: malrwahnih@ucdavis.edu