GGB Student Symposium 2013

8:00-9:00AM New Student Orientation

8:30-9:30AM Breakfast and Registration

(Genomics Lobby)

9:30-10:45AM Session I

Chair: Stephanie Coffman

9:30AM Welcome & Opening Remarks 9:45AM **Amanda Hollowell**, Sachs Lab

Variation in antibiotic resistance in natural populations of

Bradyrhizobium

10:00AM **Keira Lucas,** Raikhel Lab

Conserved microRNA miR-8 is an essential regulator of reproductive events in the fat body of the female mosquito,

Aedes aegypti

10:15AM **Bo Zhao**, Raikhel Lab

Towards improvement of genetics tools for mosquito research: Development and application of the binary Gal4/UAS system in

Aedes aegypti

10:30AM **Shaofang Li**, Chen Lab

Genome-wide profiling of Pol IV-dependent siRNA precursor

transcripts

10:45-11:00AM Coffee Break

11:00-12:00PM Session II

Chair: Caroline Sjorgen

11:00AM **Evelien Bunnik**, Le Roch Lab

Polysome Profiling reveals Translational Control of Gene Expression in the Human Malaria Parasite *Plasmodium*

falciparum

11:15AM **Stephanie Coffman**, Ding Lab

Identification of genes required for RNAi-mediated antiviral

immunity in *C. elegans*

11:30AM **Jinfeng Lu**, Ding Lab

RNA interference functions as an antiviral immunity in

mammals

11:45AM **Ilva Cabrera**. Borkovich Lab

Prediction of Gene Functions using Phenomics in a Model

Filamentous Fungus

12:00-1:30PM Lunch (HUB 355)

Session III 1:30-2:30PM

Chair: Mitchell Lucas

1:30-2:30PM **Keynote Address**

Dr. Bradley White, Entomology Department

2:30-2:45PM Coffee Break

Session IV 2:45-3:45PM

Chair: Steven Ahrendt

2:45PM Mitchell Lucas, Close Lab

> Application of Genomic Resources to Aid a Legacy of Collaborative Cowpea Breeding between UCR and Africa

Sarah Perry, Ray Lab 3:00PM

Genetic and epigenetic mechanisms underlying olfactory

learning and memory in Drosophila melanogaster.

3:15PM **Meenkashi Kagda**, Judelson Lab

Metabolic adaptations and phosphagen kinases in the plant

pathogen Phytophthora infestans

3:30PM **Jolly Shrivastava**, Judelson Lab

Building a pipeline to identify antigenic peptides specific to

Phytophthora infestans

Coffee Break 3:45-4:00PM

Session V 4:00-4:45PM

Chair: Keira Lucas

4:00PM **Steven Ahrendt**, Stajich Lab

Comparative Analysis of Putative Rhodopsins in Early

Diverging Fungal Lineages

4:15PM **Caroline Sjorgen**, Larsen Lab

Aluminum promotes root growth inhibition through an ATR-

and SOG1-dependent transcriptional response.

Gabriel Murillo, Cui Lab 4:30PM

High-Throughput Sequencing Analysis Using the REMix Base-

Caller and the GeMS SNP Caller

5:30-6:30PM Poster Session and Reception

(Genomics Lobby)

Awards and Closing Remarks 6:30PM