International Symposium on Functional Genomic Tools in Honey Bees

<u>Saturday, Aug 6, 2011</u>

112 Forest Resources Building, Penn State

8-8:30 -	Coffee, breakfast
8:30	Welcome
8:40 am	Kate Ihle, Smithsonian Tropical Research Institute, Panama
	The use of RNA-interference as a tool in large-scale behavioral studies.
9:00	Gerard Leboulle, Free University of Berlin, Germany
	The glutamatergic neurotransmission of the honey bee brain and its involvement in behavior
9:20	Ying Wang, Arizona State University, USA
	Failures of knockdown by dsRNA
9:40	Presentations from short course students (10 min/talk)
	Vanina Vergoz, University of Sydney, Australia
	Biogenic amine receptor gene expression in the ovarian tissue of the honey bee Apis mellifera
	Etya Ansalem, University of Tel Aviv, Israel
	Juvenile hormone regulation on aggression-linked differences of Vitellogenin expression in Bombus terrestris workers
	Thomas Münz, University of Würzburg, Germany
	A potential role of synapsin in the synaptic plasticity of the mushroom body calyx
10:10 am	Discussion
10:30	Coffee Break
11:00	Gro Amdam, Arizona State University/Norwegian University
	of Life Sciences, USA/Norway
	Lactobacillus kunkeei — a paratransgenic transformation system for honey bees

11:20	Michelle Flenniken (UC San Francisco, USA)
	Double-stranded RNA mediated antiviral responses in honey bees
11:40	Presentations from short course students (10 min/talk)
	Carolin Ratzka, University of Würzburg, Germany
	Does the immune system play a role in controlling the endosymbiosis of the ant Camponotus floridanus?
	Amanda van Haga, University of British Columbia, Canada
	Characterization and validation of host-pathogen interactions between microsporidia (Nosema spp.) and honey bees (Apis mellifera L.)
	Arian Avalos, University of Puerto Rico, USA
	Memory and endonucleases: A look at the role of fen-1 in long term memory potentiation
12:10	Discussion
12:30	Lunch
1:30 pm	Christina Grozinger, Penn State University, USA
	Behavioral and evolutionary genomics of pheromone communication
1:50 pm -	Ryszard Maleszka, Australian National University, Australia
	Five years of honey bee genomics: what we now know, what we do not know, and what we can do?
2:10 am -	Zilá Simões, University of São Paolo, Brazil
	Using next generation tools for discovery of developmentally regulated genes
2:30 am	Presentations from short course students (10 min/talk)
	Aline Silva, University of São Paolo, Brazil
	Molecular determinants in worker Apis mellifera development
	Karl Glastad, Georgia Tech University, USA
	Utilizing RNAi to explore DNA methylation and mRNA splicing
	Kathe Munke, University of Aarhus, Denmark
	RNAi induced alterations of methylation patterns in Honey Bees

3:20	Coffee Break
3:40	Yehuda Ben-Shahar, Washington University, USA
	The 'social' fly: using comparative genetics to understand sociality
4:00	Ehab Abouheif, McGill University, Canada
	Ant polyphenisms
4:20	Presentations from short course students (10 min/talk) Rajee Rajakumar, McGill University, Canada Using RNAi to understand the potential role of gene network interruption points underlying wing polyphenism in ants.
	Hagai Shpigler, Hebrew University of Jerusalem, Israel
	Using RNAi technology to explore the function of Krüppel- homolog 1 in the social organization of honey bee and bumble bee societies
	Carolina Santos, University of São Paolo, Brazil
	Differentially expressed genes in hind leg development of honey bee larvae
4:50	Discussion

5:20 Final comments

Sunday, August 7

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Discussions 9-12 pm (with break at 10:30)

Lunch 12-1

Wrap up session 1-2

Topics for discussion:

- 1. Status of the RNAi field and where we are in it
- 2. What are our options and strategies for progress?
- 3. What critical experiments can and should be done?
- 4. Who will do these experiments?
- 5. How can results be communicated effectively (prepublication?)?,