Symposium and Workshop on New Methods for Phylogenomics and Metagenomics

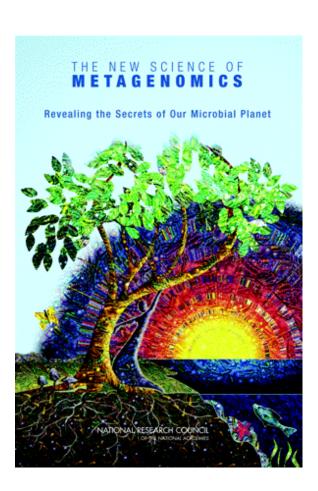
February 16 and 17, 2013

Tandy Warnow
National Science Foundation
University of Texas at Austin

Computational Phylogenomics and Metagenomics







Symposium talks

- Phylogeny/alignment estimation
- Phylogenomics
- Phylogenetic placement
- Metagenomics
- 1KP (Thousand Transcriptome Project)

NSF DEB 0733029

- Title: <u>Large-Scale Simultaneous Multiple Alignment and Phylogeny</u> Estimation
- Collaboration between
 - Texas (Pingali, Warnow, and Linder), lead institution
 - Kansas (Holder),
 - Nebraska (Moriyama),
 - Georgia (Leebens-Mack)
 - Subawards: Smithsonian (Braun), NJIT (Roshan), and Penn State University (Miller)
- Project webpage

http://www.cs.utexas.edu/users/tandy/ATOL-MSA.html

Previous Workshops

- NESCENT 2011
- Smithsonian 2012

We will seek funding for another workshop for 2014.

Saturday Schedule

- 8:00-8:45 AM Breakfast in NHB 1.720 pick up name tags!
- 9:00 12:00 Talks in ACES 2.302 (Avaya Auditorium)
- 12:15 -1:15 Lunch in NHB 1.720
- 1:15 3:00 Talks
- 3:00 3:30 Coffee Break
- 3:30 5:00 Talks
- 5:00 5:30 Discussion

Tutorials (Sunday)

- All tutorials in the Avaya Auditorium are open (SATé, Phylonet, BUCKy, and *BEAST). Other tutorials (held in ACES 2.402 and PAT 141-SA) are limited in number of seats.
- BAli-Phy is closed.
- If you are interested in attending the other tutorials and did not already register, please contact me by 1 PM today by email (<u>tandy@cs.utexas.edu</u>) to get on the waitlist.
- Today: Please download, install, and test the tutorial software.
 Some tutorials (e.g., SATé and SEPP) also have written tutorials that can be downloaded in advance.
- Breakfast on Sunday is at 7:30, and the tutorials start at 8:30.
 Please be in your seats a few minutes ahead of time, as the schedule is tight!

Logistics

- Laurie Alvarez (CCBB Senior Administrator), Nicole Elmber, Anna Akhtar, or Tandy Warnow for assistance. Email: tandy@cs.utexas.edu
- Please wear your name tag while at the conference
- NO FOOD OR DRINKS (other than water) allowed in the meeting rooms
- "Bio-breaks" as needed take two right turns as you exit the auditorium.

 Schedule is tight - please be in your seat a few minutes ahead of the start time

Meeting Objectives

- Stimulate research into new methods
- Hands-on training in new methods
- Obtain feedback about new methods
- Develop collaborations between methods developers and biologists

Conference Support From

NSF DEB-0733029 (\$100K)

 The Center for Computational Biology and Bioinformatics (CCBB) at UT-Austin (Laurie Alvarez)