**Application for Travel Award to the**

**2015 Phylogenomics Symposium and Software School**

**Please print legibly (or type), and send along with your two-page CV, to Tandy Warnow (**[**warnow@illinois.edu**](mailto:warnow@illinois.edu)**) with subject “Phylogenomics Symposium”. If you are a student or postdoc, also ask your advisor to send a recommendation letter to the same email address. Applications received by April 1 will receive first consideration, and travel awards will be announced by April 15, 2015. Travel awards are only awarded to participants who plan to attend both days – the symposium and tutorials.**

**Name: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**

**University: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**

**Department: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**

**Homepage: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**

**Email address: \_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_\_**

**Current position (please circle one):**

**Faculty**

**Staff Scientist**

**Postdoctoral fellow**

**PhD student**

**MS student**

**Undergraduate student**

**Other (explain)**

**1. Experience with phylogenetic estimation methods (circle one)**

**Expert**

**Experienced but not expert**

**Have done this a few times**

**Just starting**

**2. Experience with multiple sequence alignment methods (circle one)**

**Expert**

**Experienced but not expert**

**Have done this a few times**

**Just starting**

**3. Experience with coalescent-based species tree estimation (circle one)**

**Expert**

**Experienced but not expert**

**Have done this a few times**

**Just starting**

**4. If you are not a faculty member, please give the name and email address of your faculty advisor. (If this is not applicable, explain.)**

**5. Which multiple sequence alignment methods have you used, and which ones do you prefer (and why)?**

**6. Which phylogeny estimation methods have you used, and which ones do you prefer (and why)?**

**7. Which coalescent-based species tree estimation methods have you used, and which ones do you prefer (and why)?**

**8. Briefly describe your research, how it relates to multiple sequence alignment and phylogeny estimation, and why this summer school is useful to you.**

**9. Which of the following tutorial topics do you wish to attend? (Rank them by importance to your research, with 1 being the most important and**

**\_\_\_\_\_\_ Multiple sequence alignment methods (two tutorials - PASTA and UPP)**

**\_\_\_\_\_\_\_\_ Coalescent-based species tree estimation (two tutorials – ASTRAL and SVDquartets)**

**\_\_\_\_\_\_\_\_ Phylogenetic network estimation (one tutorial – Phylonet)**

**\_\_\_\_\_\_\_\_ Metagenomic taxon identification (one tutorial – TIPP)**

**10. Please identify the laptop you will bring (and the operating system) – linux machines and Macbooks running OS X are preferred compared to PCs running Windows!**

**11. If you will need financial support, please indicate the total amount that you would request, and the basis on which you calculate this. If you have other sources of funding, please indicate this.**