

# Home

## Welcome to WiNGS

WiNGS is a symposium and workshop introducing software tools and pipelines for processing, visualization, exploration, and analysis of biological data sets. The goal of WiNGS is to help life scientists learn to use bioinformatics tools and pipelines in their research.

## About this Wiki

This collaborative wiki page contains materials from the 2013 and 2014 WiNGS workshops, hosted by the [Department of Bioinformatics and Genomics](#) in the [College of Computing and Informatics](#) at [UNC Charlotte](#).

## Credits

WiNGS receives support from the National Science Foundation grant number [1238051](#) titled **Genomic Approaches to Understanding Cytokinin Signaling and Function in Rice** led by Principal Investigator [Joe Kieber](#) (UNC Chapel Hill) and co-PIs [Ann Loraine](#) (UNC Charlotte) and [Eric Schaller](#) (Dartmouth University).

This wiki is provided by the [iPlant Collaborative](#). If you have questions about how to edit the wiki or about other iPlant resources, please feel free to contact [iPlant Collaborative Support](#).