

NEXTFLOW & CONTAINERS WORKSHOP

January 16-18 9am - 12pm

Sign up to learn the fundamentals of Nextflow and containers:

- *Develop flexible bioinformatics pipelines using Nextflow for reproducible science*
- *Find, deploy, and build custom Docker images*
- *Run Singularity containers in HPC environment*

Experience with command-line highly recommended

github.com/statonlab/NF_workshop

WHERE

160 Plant Biotechnology
Knoxville, TN

COST

\$100

SIGN-UP

To sign up for the waiting list, please contact
rkuster@utk.edu

