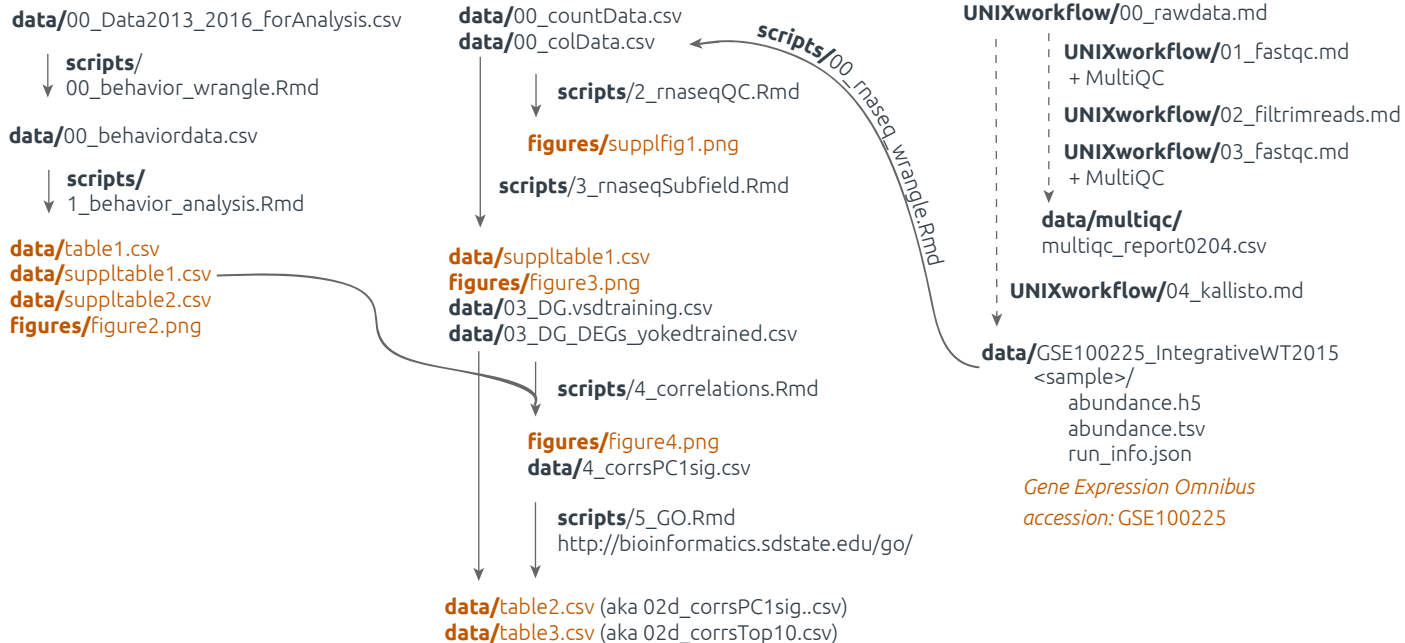


Bioinformatic workflow for Harris *et. al* 2020

<https://github.com/raynamharris/IntegrativeProjectWT2015/>

R + GitHub

Texas Advanced Computing Center



Visit <https://github.com/raynamharris/IntegrativeProjectWT2015> and click "launch binder" to explore the R workflow in the cloud with an myBinder.

Notes: R scripts beginning with "00" are all about data wrangling. Focus on scripts beginning with 1-5 to reproduce the manuscript.

The UNIX workflow is not executable; rather, it is a narrative containing the UNIX commands that were executed.

Outputs for the manuscript are highlighted in burnt orange.