



Identify

BagIt

Completeness



Access



RO-Crate

Describe and Relate



BioCompute
Objects

Explain

Execute



CWL

Exemplify

results/

- fastqc
- genome
- igv/broadPeak
- multiqc/broadPeak
- pipeline_info
- trim_galore

Attribution



License



docker

BIOCONDA

Software Packages