Raw Data

Results

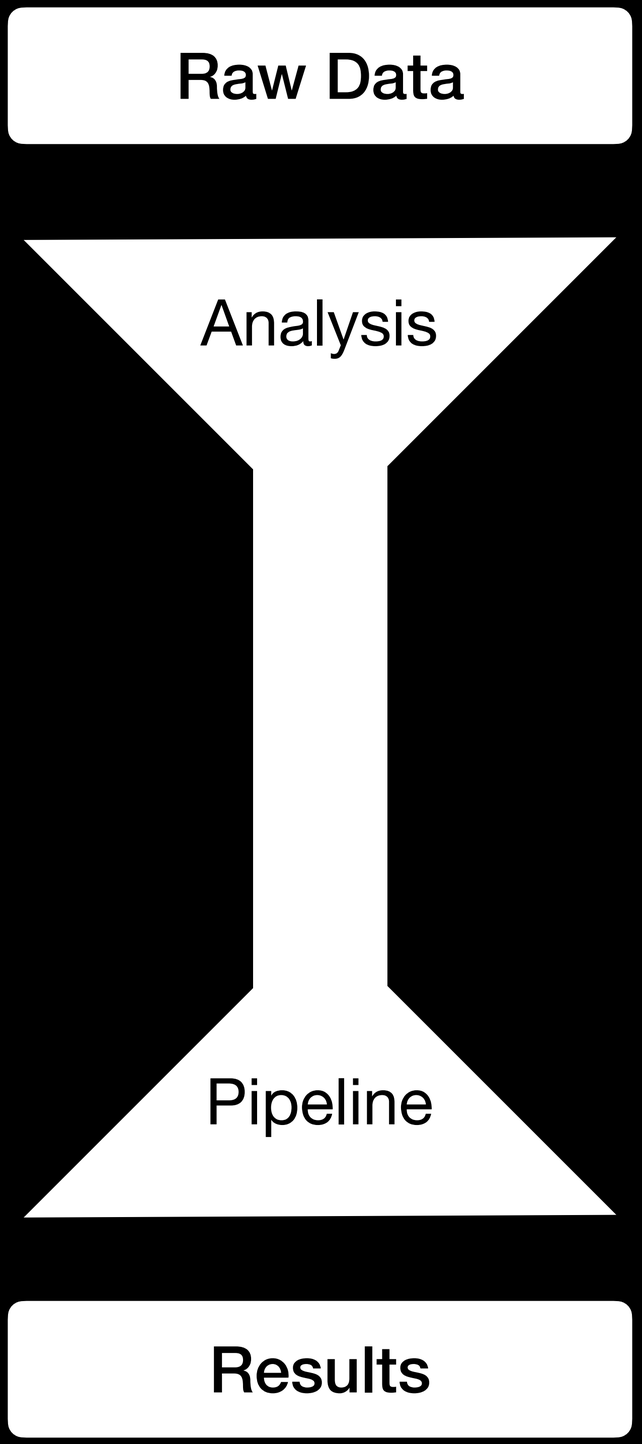
Analysis

Pipeline

Dosage Compensation Pipeline

Input:

- RNA-Seq Data
- Newick Tree with divergence estimate in MYA
- At least one species with a reference transcriptome annotation.



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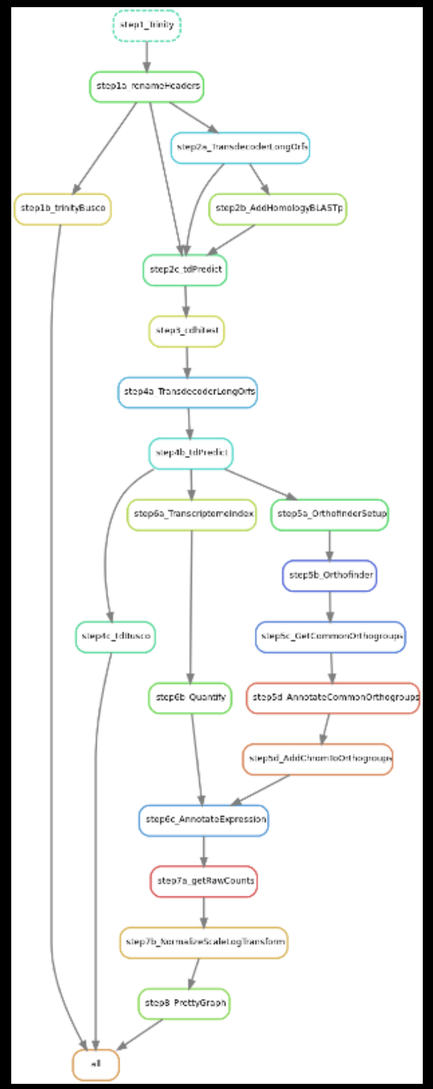

Results