



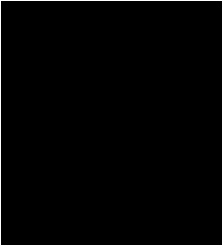
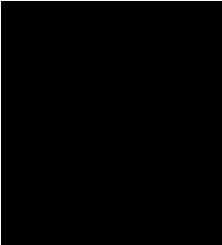
T. brev

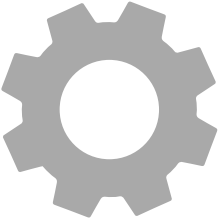


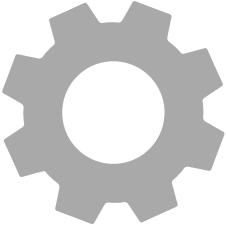
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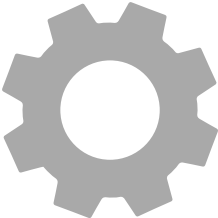


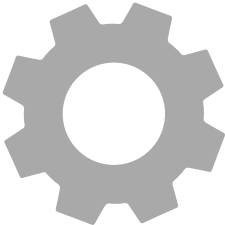
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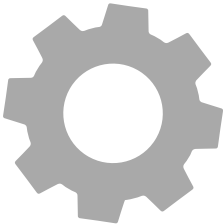


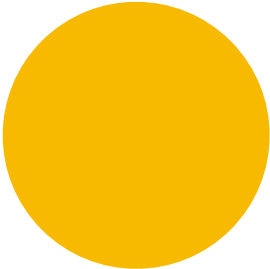


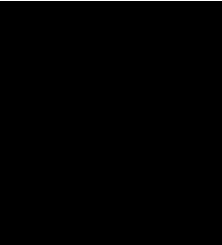
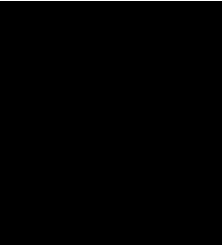


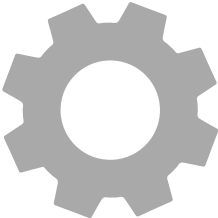


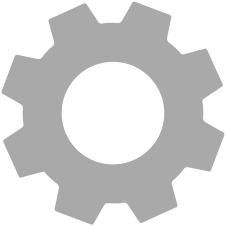


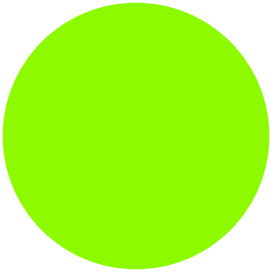


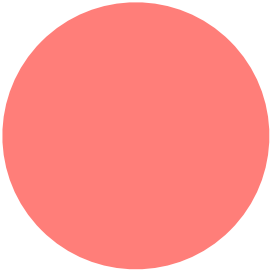


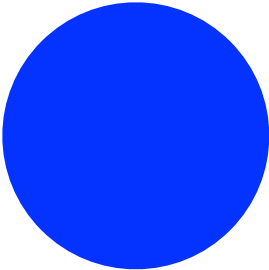


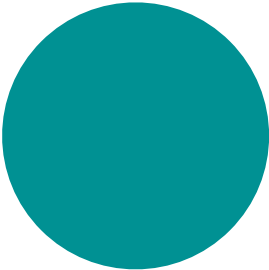


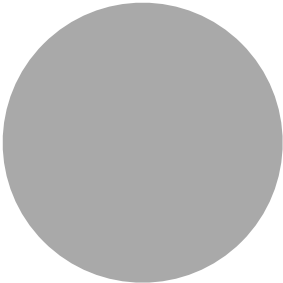


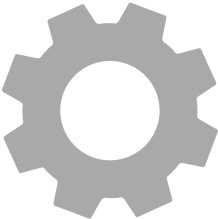


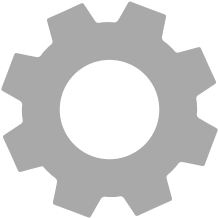












Progress

Extracted
RNA

de novo Transcriptome Assembly

RNA-Seq

T. frem

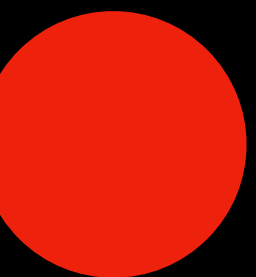
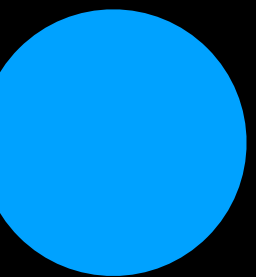
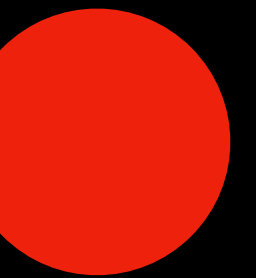
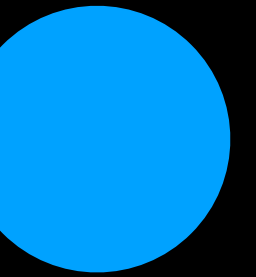
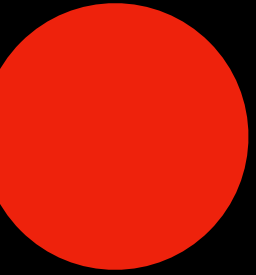
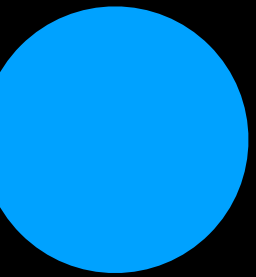
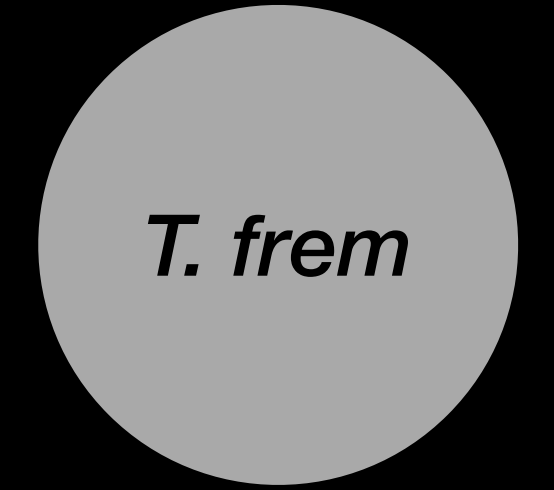
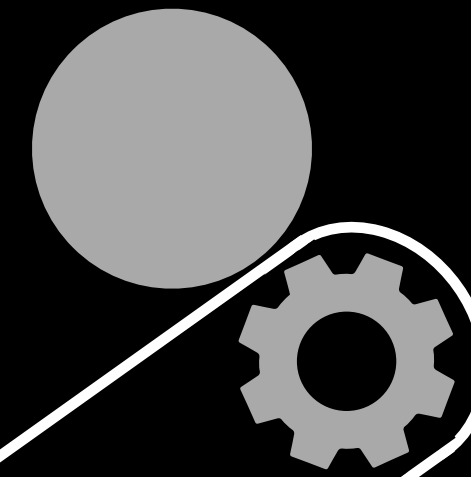
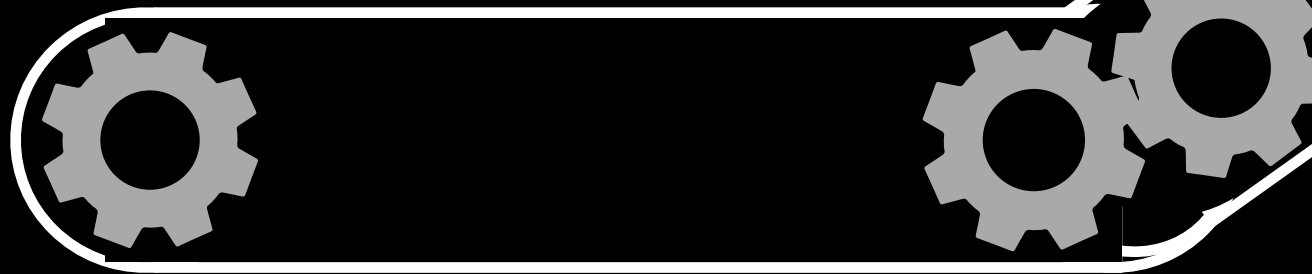
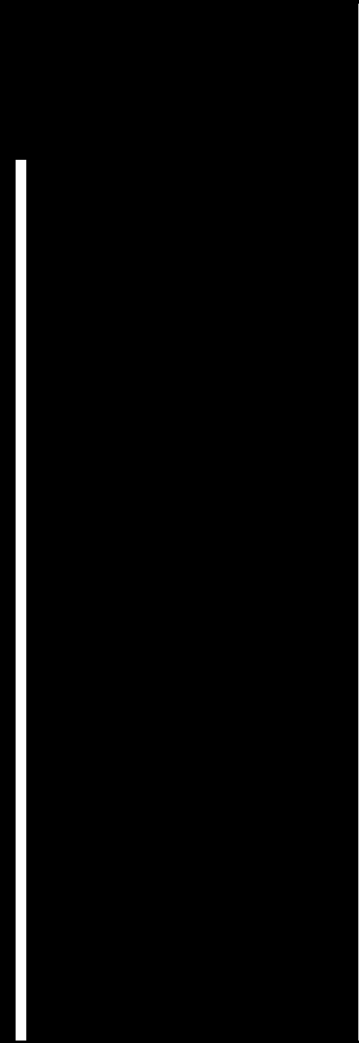
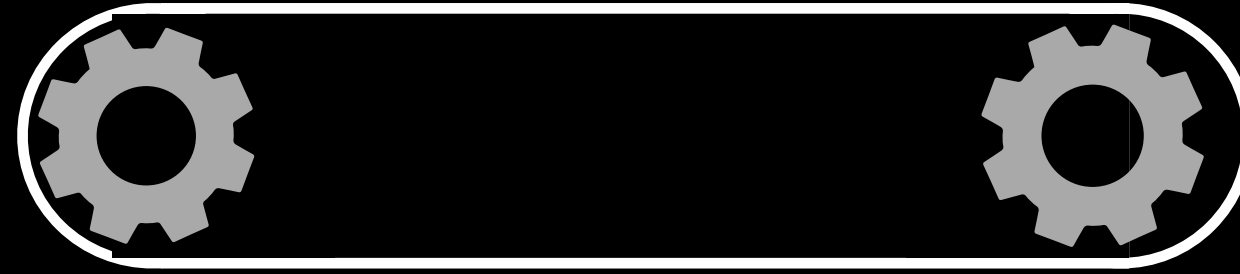
T. brev

G. corn

FastQC and MultiQC

Expression Quantification
and Normalization

TBLASTX
and
Differential
Expression
Analysis



Acknowledgements

- Phi Sigma
- Demuth Lab
 - Jeff
 - Maya
 - Catherine
 - Beena
 - Emily
 - Kayla
 - Kimmy
- Fujita Lab members
- Betrán Lab members
- Castoe Lab members
- Friends and family in LS, SIER and back home.