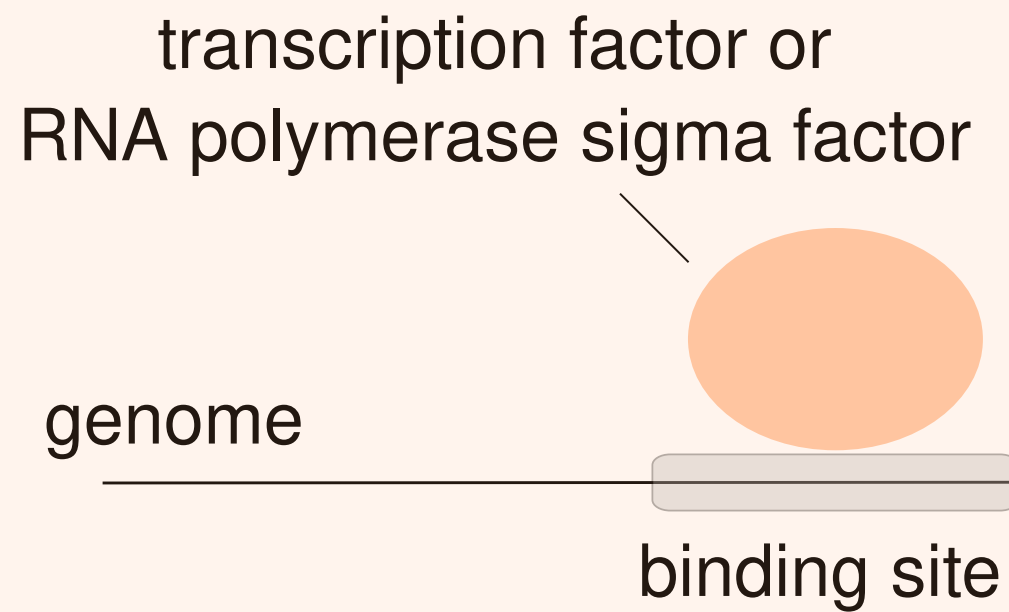
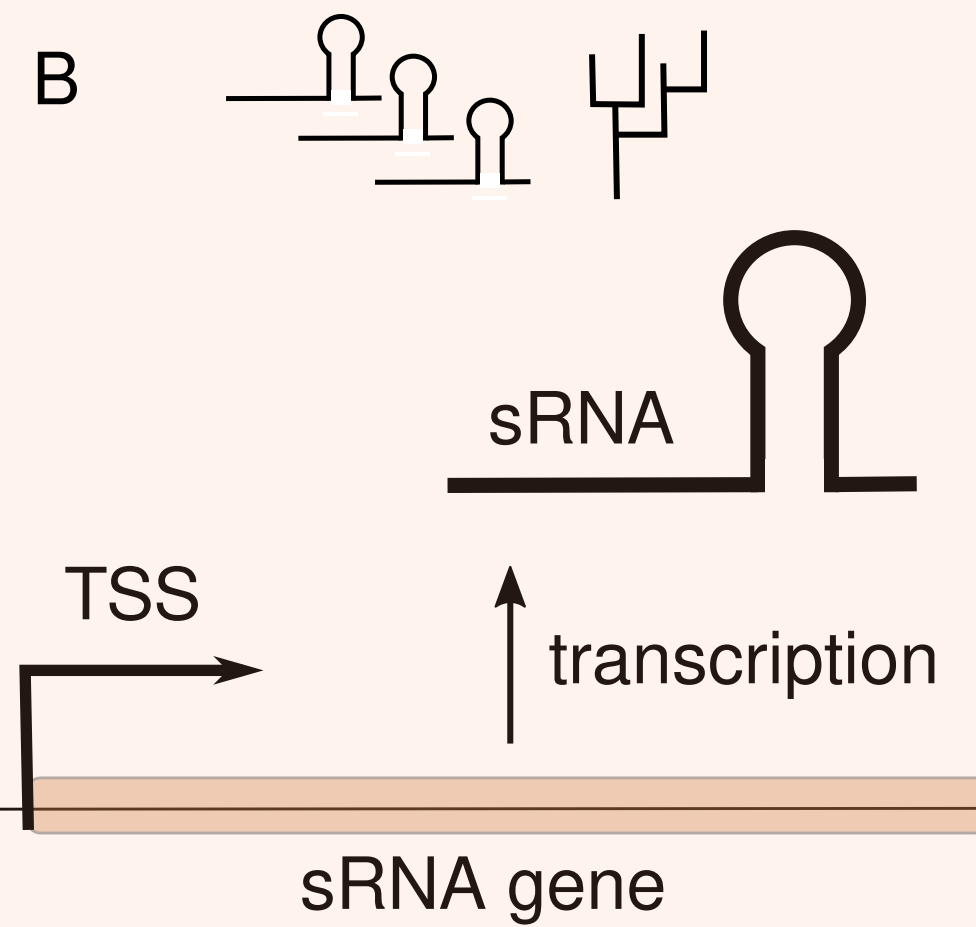


A



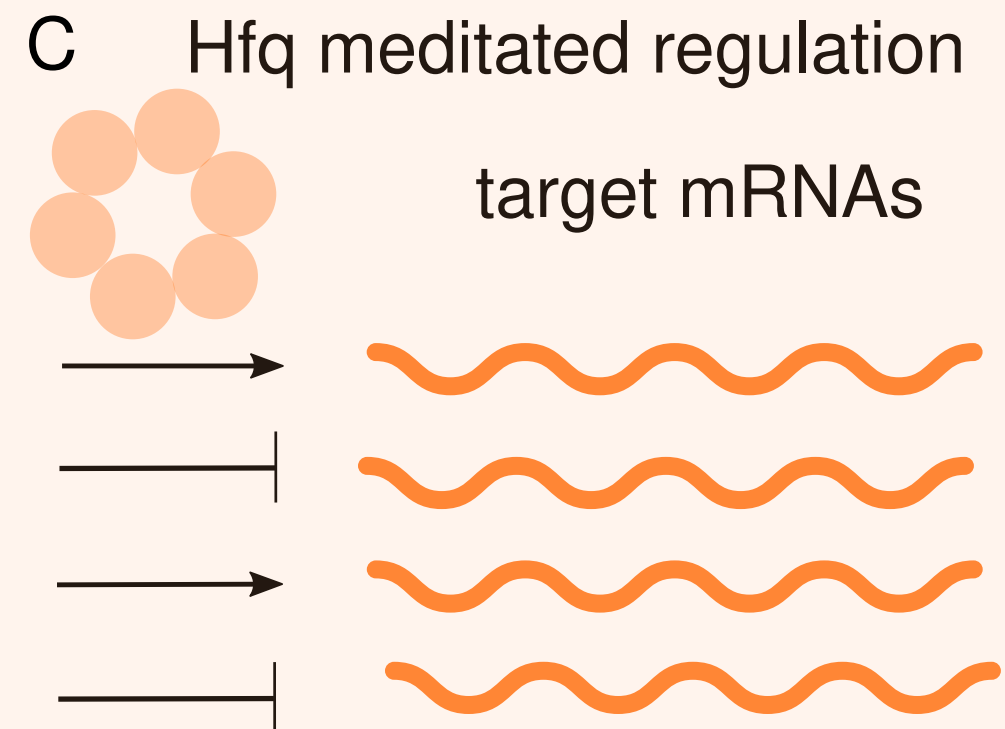
Transcription factor and
sigma factor binding site
prediction

B



sRNA annotation based
on dRNA-Seq and
RNA-Seq

C

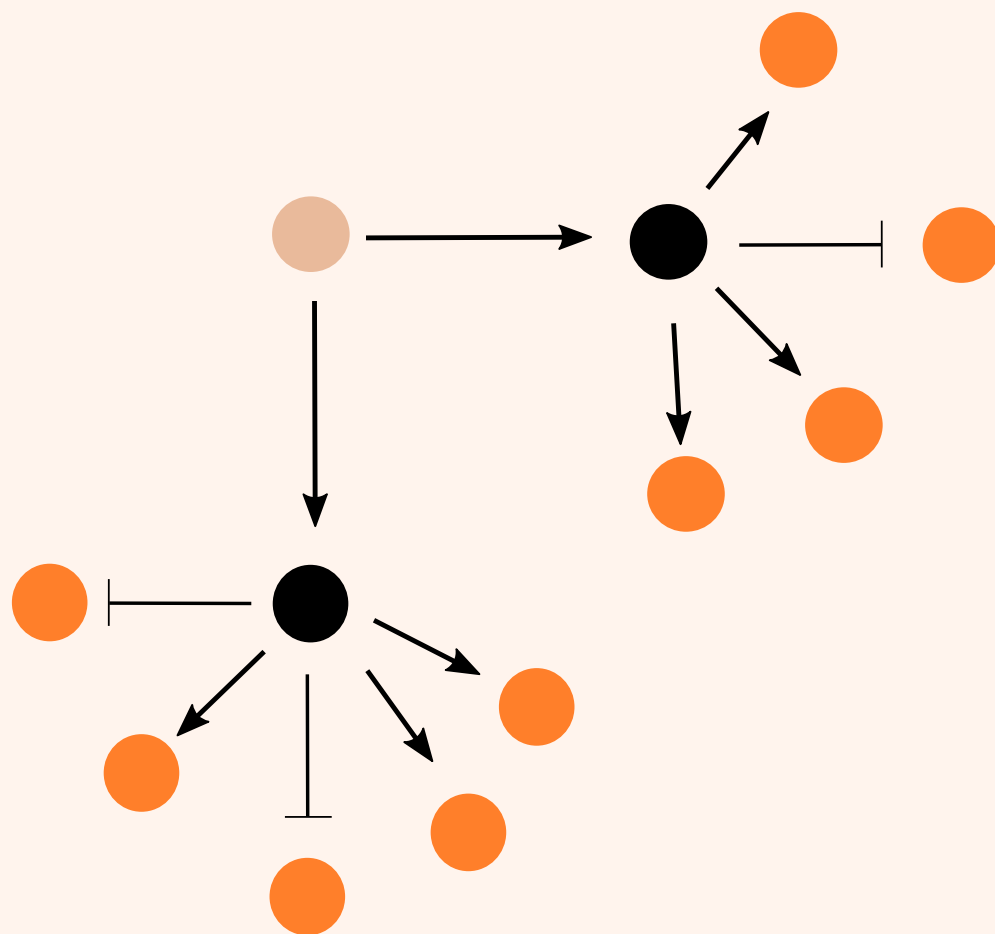


Identification of
sRNA-mRNA interactions
based on RIL-Seq

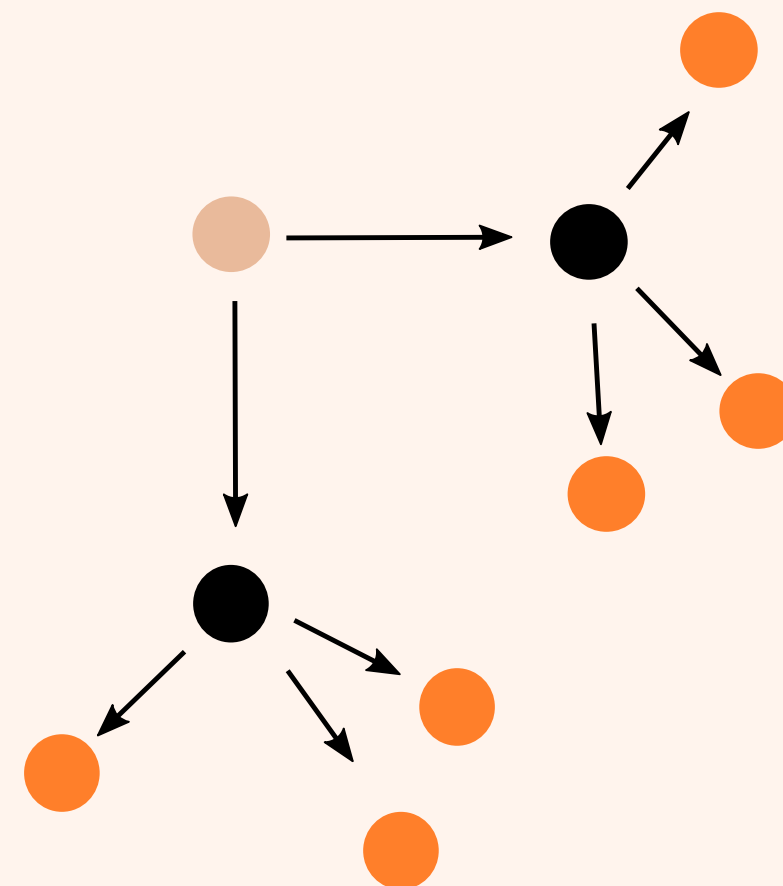
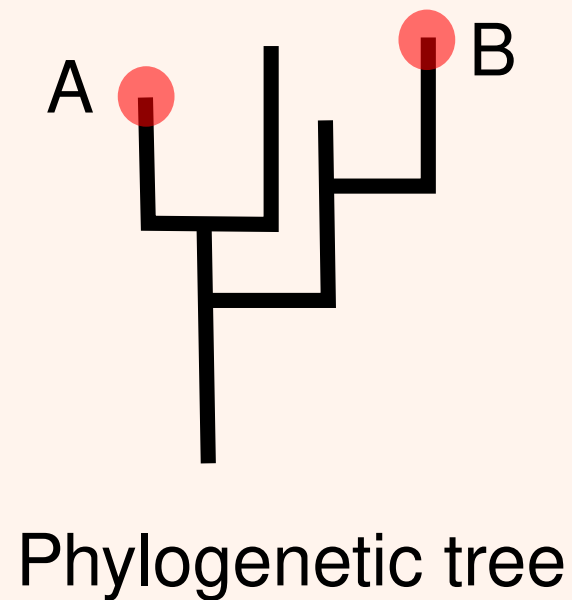


Network reconstruction

D



Regulatory network species A



Regulatory network species B

Comparative analysis of sRNA regulation networks between species