# NEEDLE for gene discovery in non-model plants

#### Question



## Input data

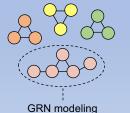
Dynamic transcriptome dataset



6 – 30 samples across variables (e.g., genotypes, tissues, or time points)

### Prediction

Coexpression network modeling





#### Validation

Promoter scanning



Transient reporter assay

