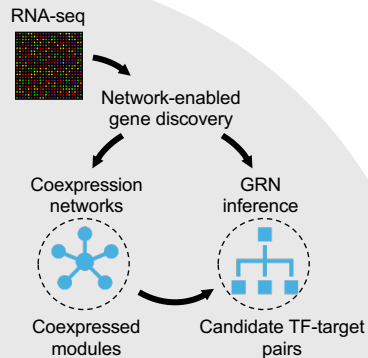
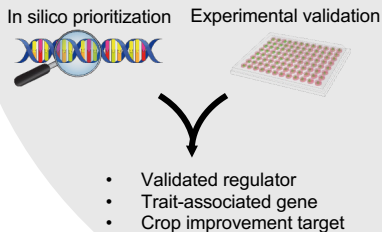


1. Biological context and regulatory challenge



2. Transcriptome-based network inference

4. Validation and gene discovery



3. Multi-omics refinement of TF–target prediction

