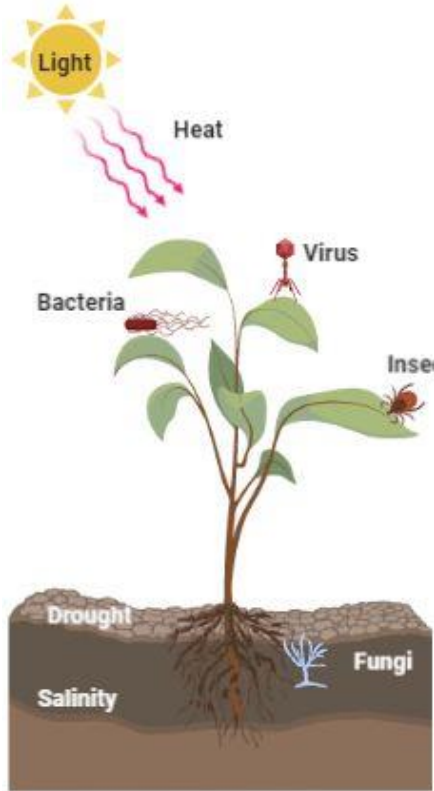


Transcriptome profiling of combined stress data

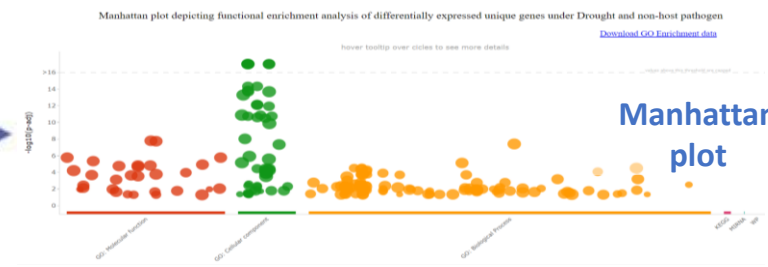
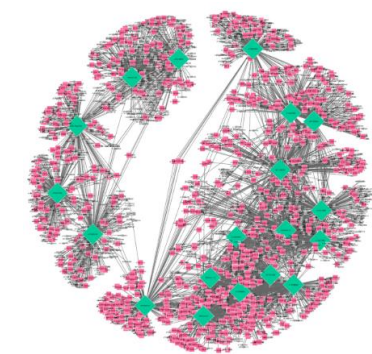
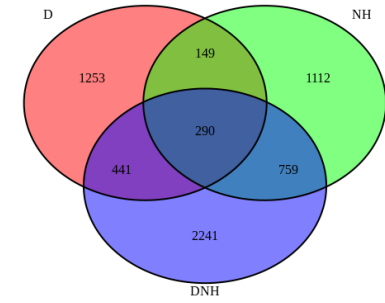
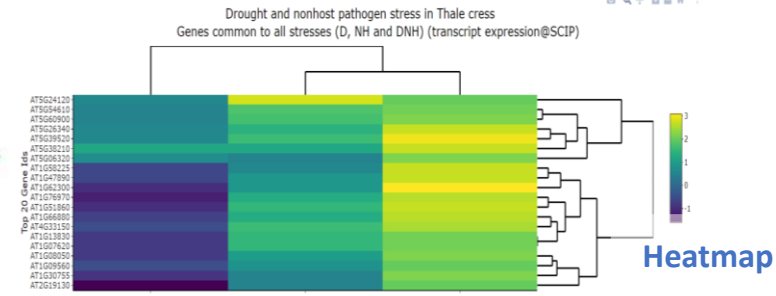


Combined Stress
 Abiotic-Abiotic
 Abiotic-Biotic
 Biotic-Biotic

Microarray Analysis

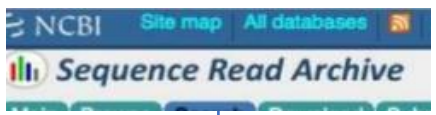
RNA Seq Analysis

Differentially Expressed Genes
 (4 Common categories and 3 unique categories)



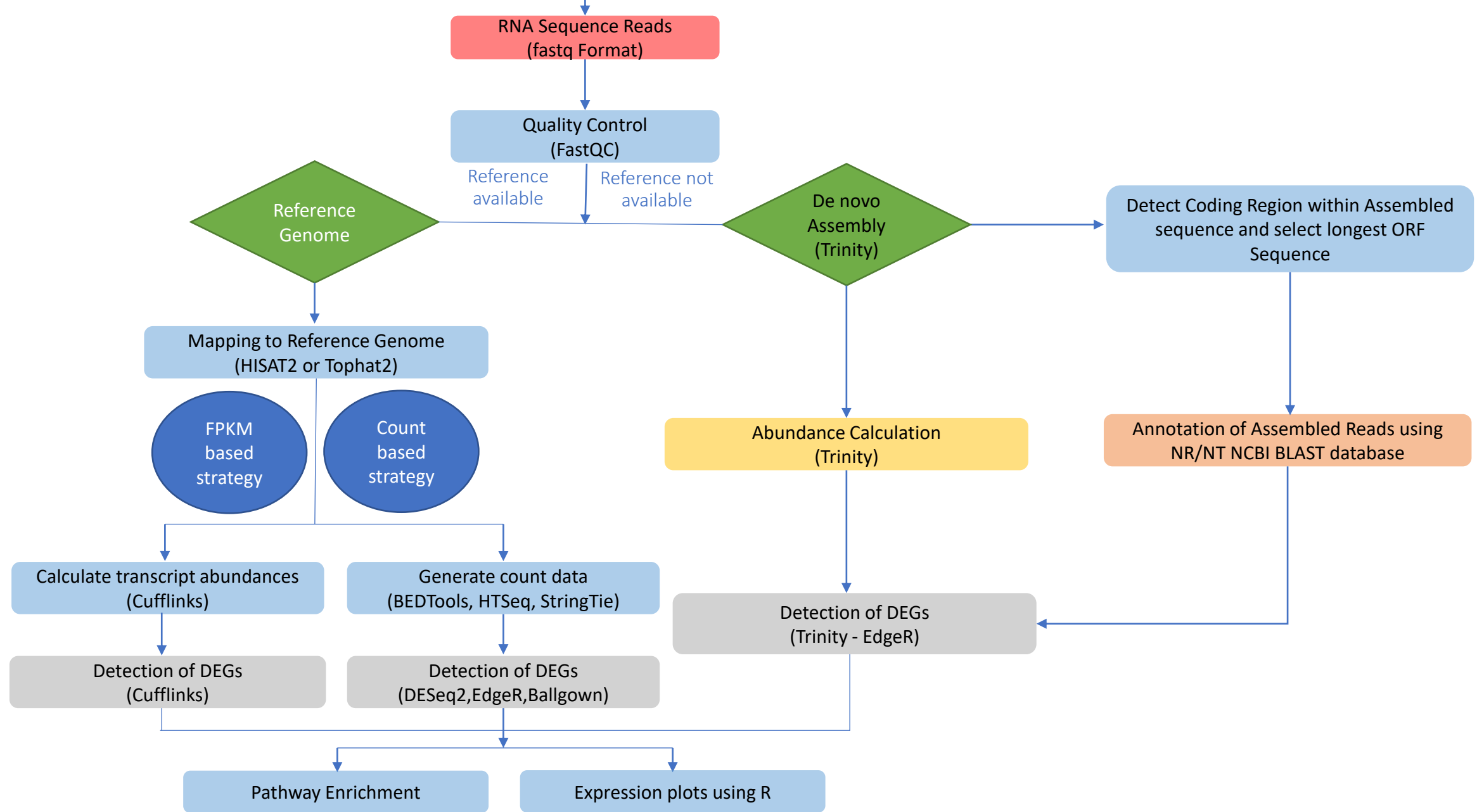
Data Collection

RNA Seq combined stress data analysis pipeline



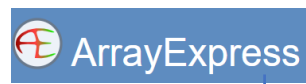
Preprocessing

Analysis



Data
Collection

Microarray combined stress data analysis pipeline



Preprocessing

Analysis

