Tool version:

Stringtie	*2.2.1*
htseq-count	*1.99.2*
Ballgown	*2.40.0*
edgeR	*4.6.2*
DESeq2	*1.48.1*
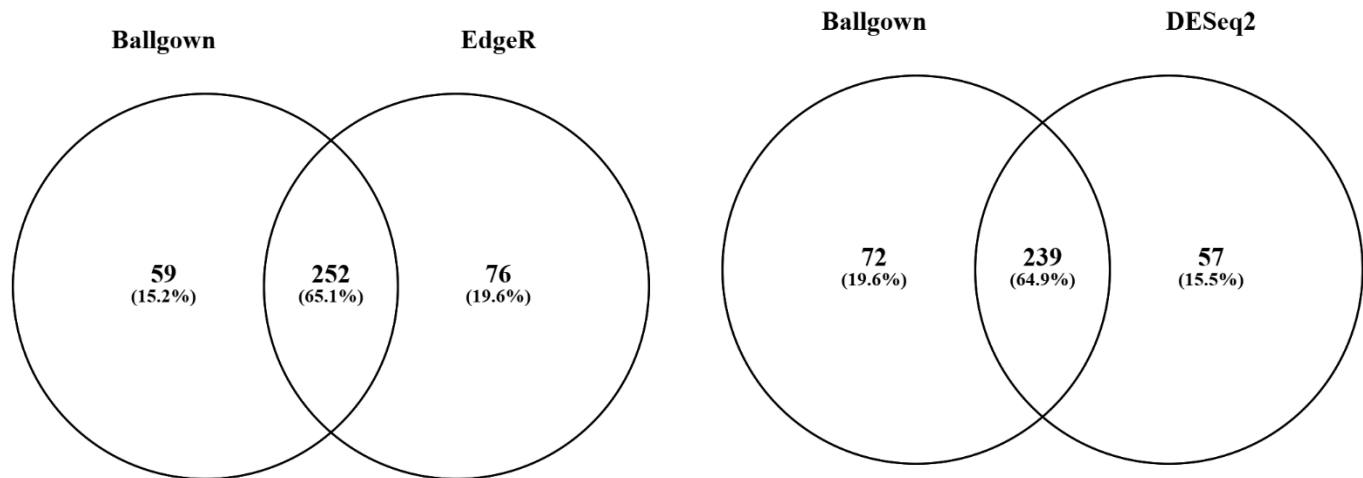
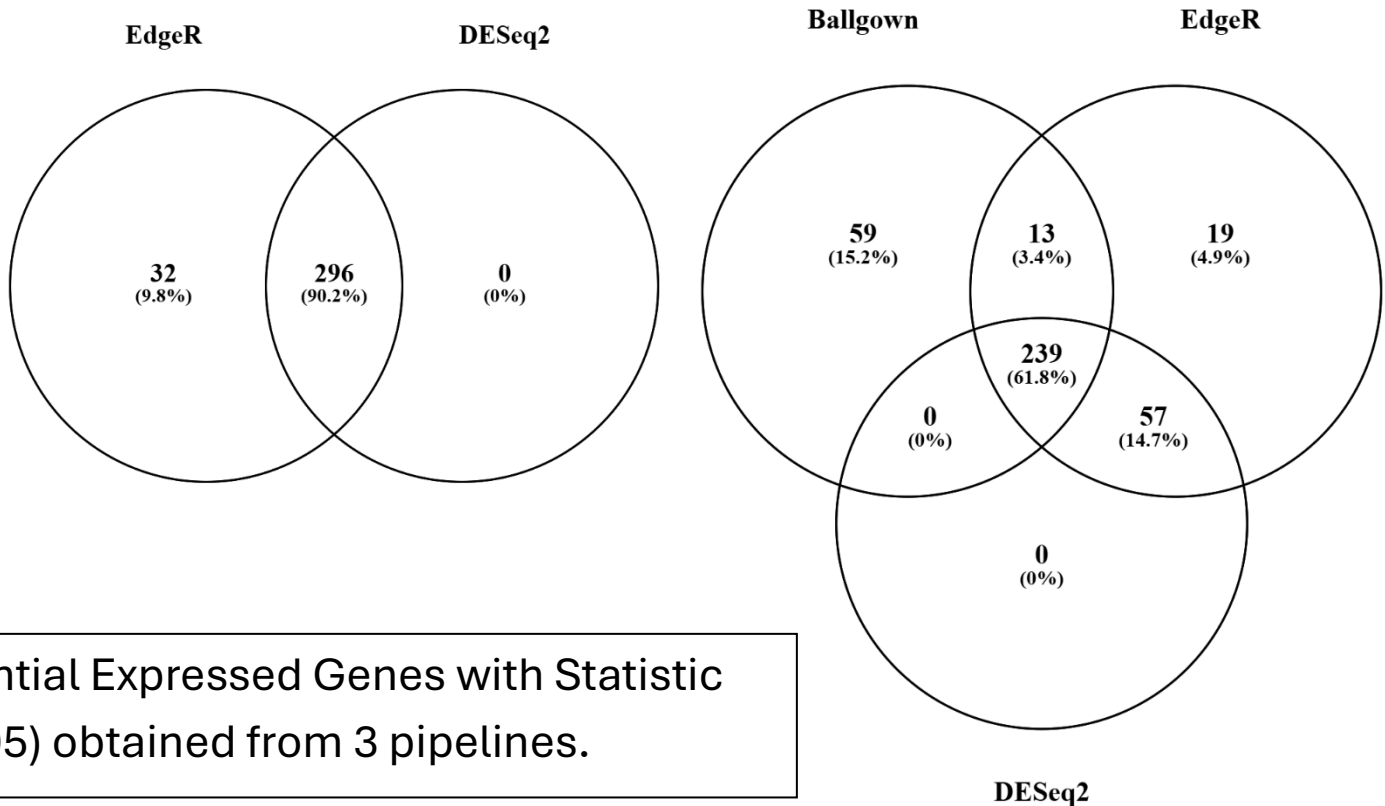Pipelines used:

Stringtie + ballgown

→ 311 significant DEG

htseq-count + edgeR

→ 328 significant DEG

htseq-count + DESeq2

→ 296 significant DEG

Overlapping Differential Expressed Genes with Statistic Significance (p < 0.05) obtained from 3 pipelines.