

Corresponding author(s): \_\_\_\_\_

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Illumina Hi-seq 2500 sequencer and Illumina Hi-seq 4000 platform

FastQC, Trimmomatic v.0.36, STAR aligner v.2.6.0a, R package edgeR v.3.22.5, AgriGO v2.0, ggplot2, Bowtie2, MACS2, MEME

*Provide your data availability statement here.*

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