

Supplementary Fig. 1

Transcriptome analysis of apical meristem enriched bud samples for size dependent flowering commitment in *Crocus sativus* reveal role of sugar and auxin signalling

Anjali Chaudhary^{1,2}, Kunal Singh^{1,2}

¹CSIR-Institute of Himalayan Bioresource Technology, Palampur-176061, India

²Academy of Scientific and Innovative Research (AcSIR), Ghaziabad-201002, India

*Correspondence to: kunal@ihbt.res.in

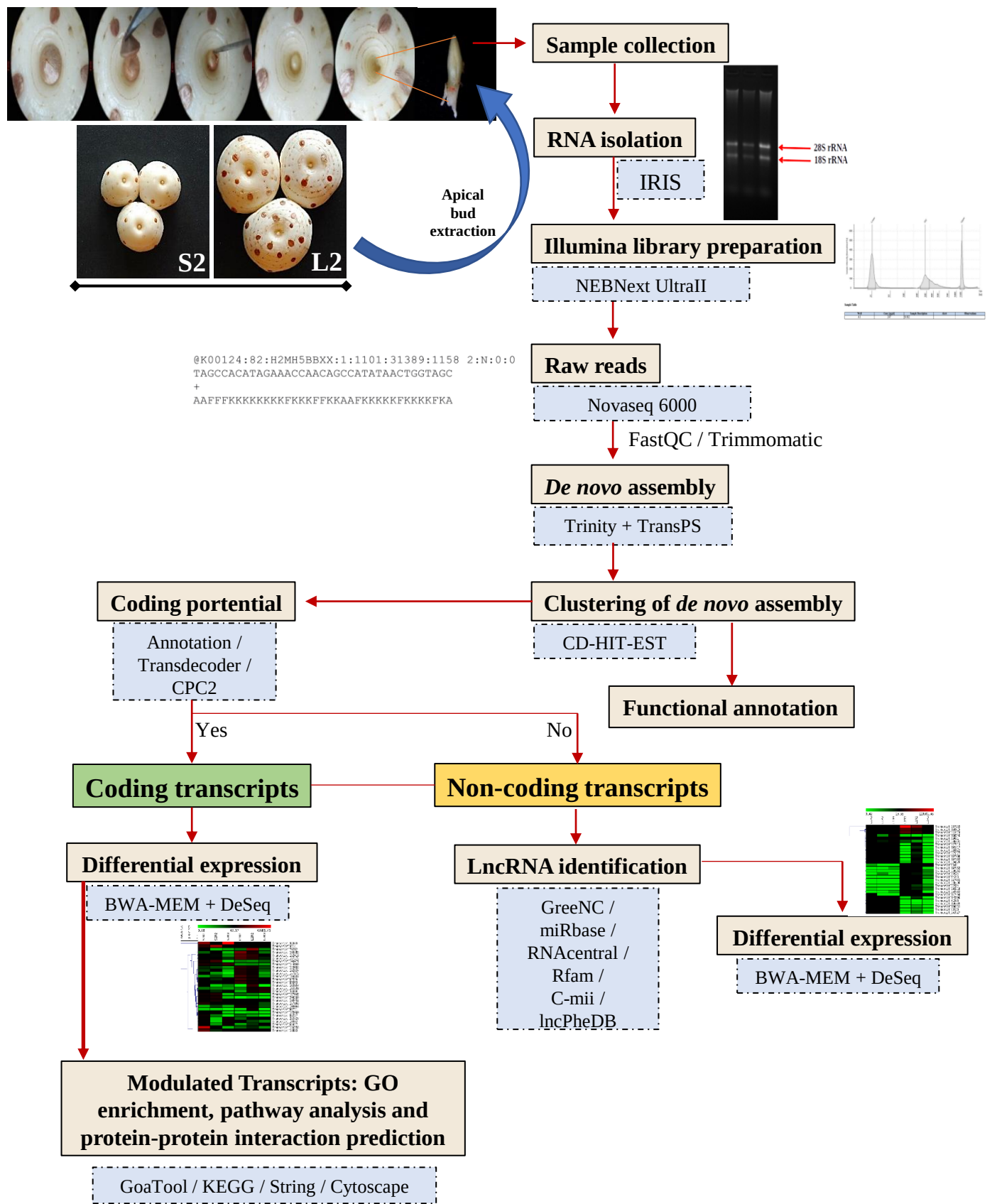


Fig. 2 Flow chart depicting the pipeline used for RNA-Seq data analysis