

Supplementary Materials: Generation and Characterisation of a Reference Transcriptome for Lentil (*Lens culinaris* Medik.)

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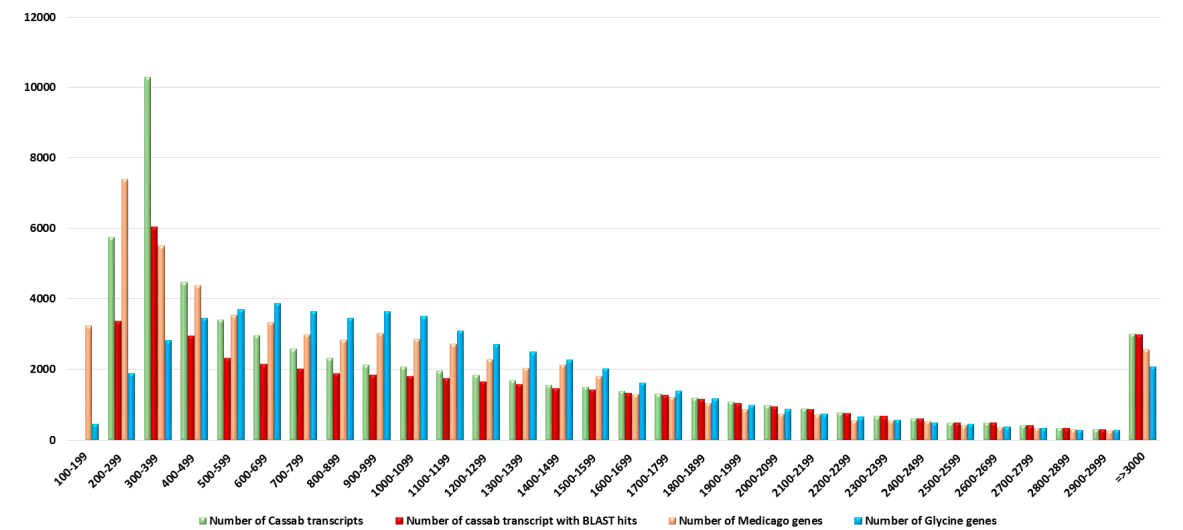


Figure S1. Length distribution of transcripts/genes—Cassab reference transcripts, Cassab reference transcripts with Nr BLAST hits, Medicago genes and Glycine genes.

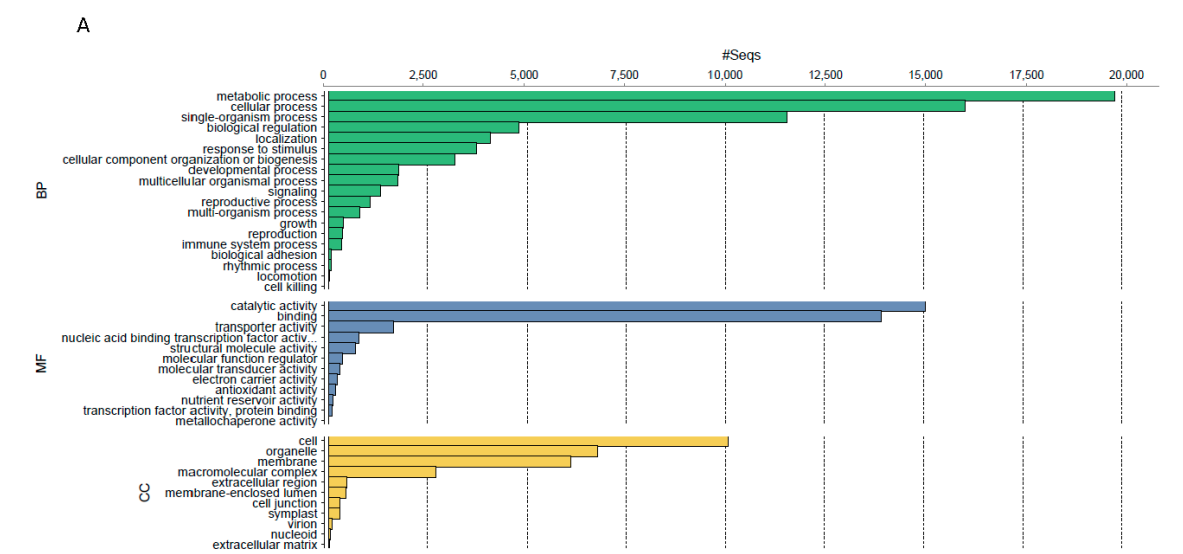
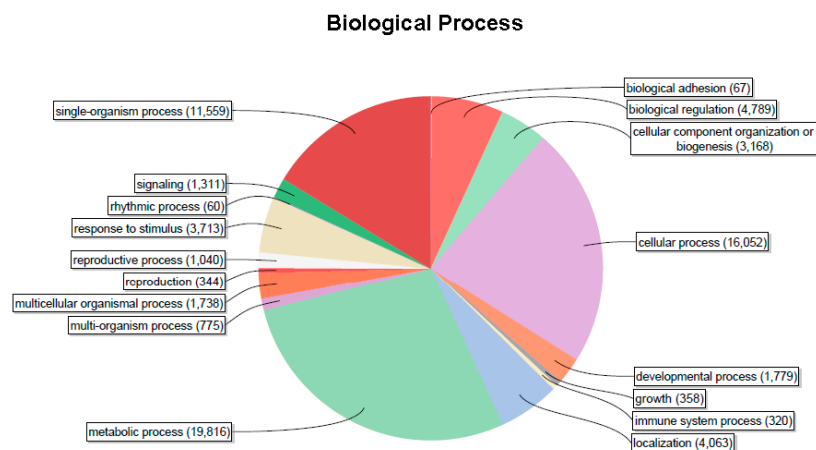
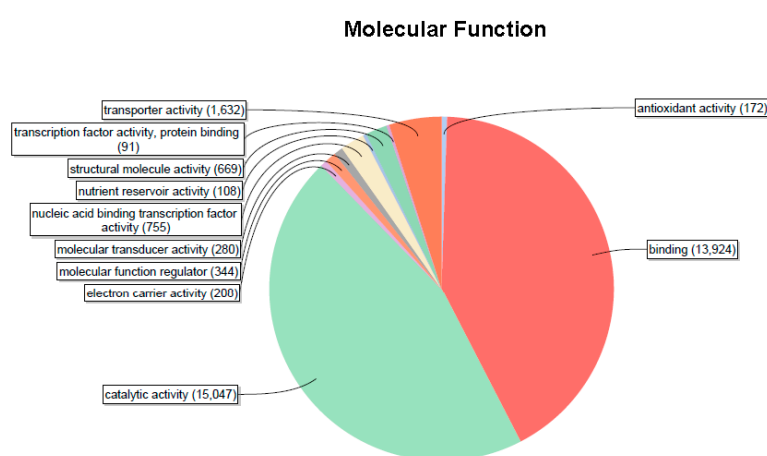


Figure S2. Cont.

B



C



D

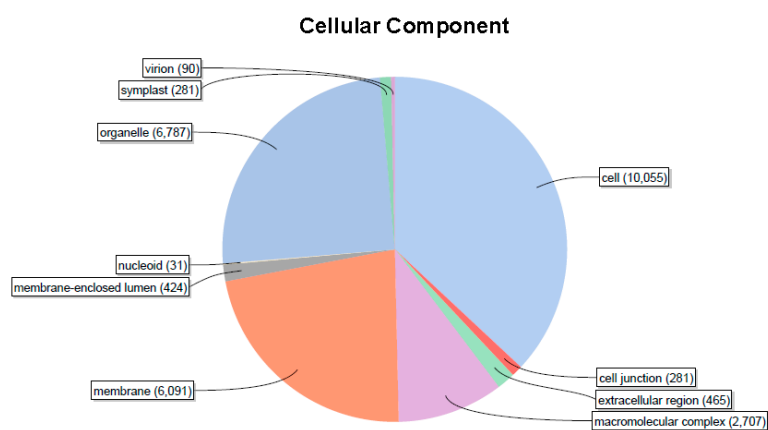


Figure S2. (A) Distribution of the annotated transcripts under gene ontology categories such as biological process (BP), molecular function (MF) and cellular component (CC); (B) Distribution of the annotated transcripts under gene ontology category–biological process (BP); (C) Distribution of the annotated transcripts under gene ontology category–molecular function (MF); (D) Distribution of the annotated transcripts under gene ontology category–cellular component (CC).