

Additional file 7:

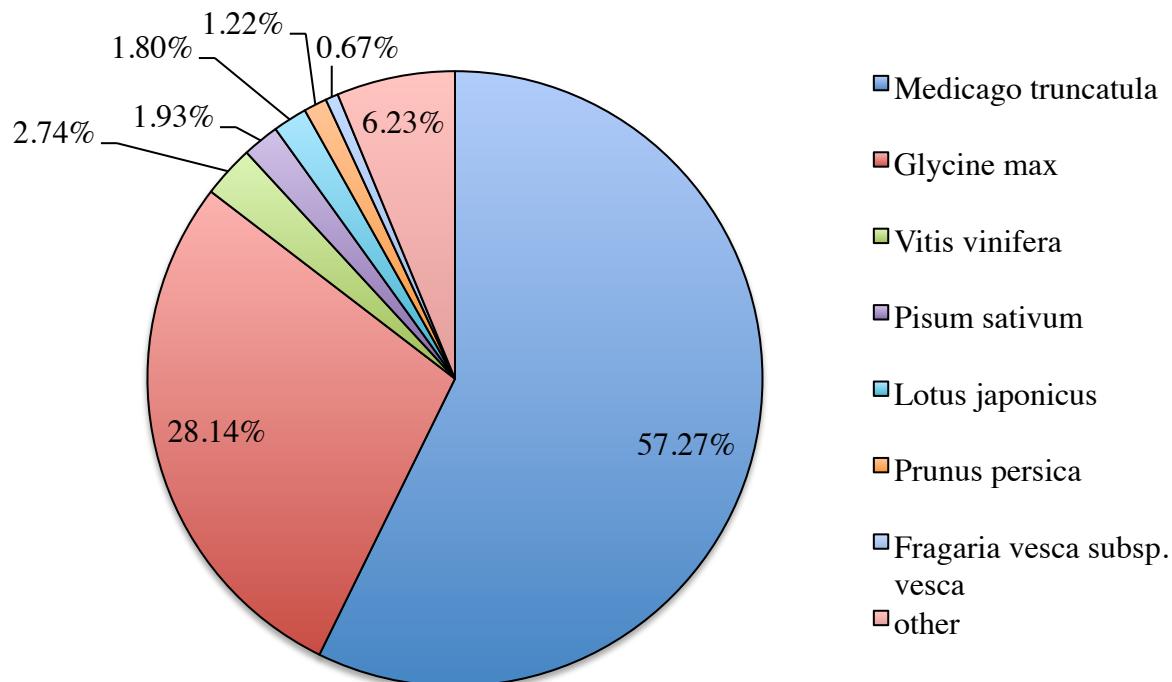


Figure S3. Species distribution of genes with highest sequence similarity (lowest BLAST E value) to unigenes in the reference transcriptome library of *trans*-differentiating adaxial epidermal cells of cultured *V. faba* cotyledons. Data obtained by aligning unigenes from the reference library with NCBI nr, Swissprot, KEGG, and COG databases using BlastX.