

Additional file 8: GO functional enrichment analysis for predicted target genes of miRNAs in four tissues respectively with comparisons to total foxtail millet genes

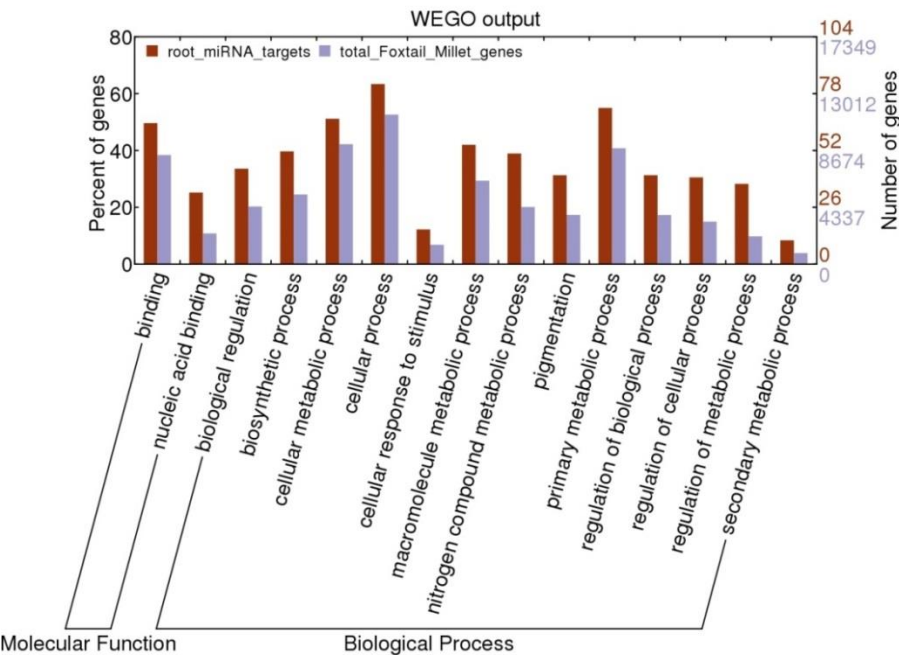
(A) GO functional enrichment analysis for predicted target genes of miRNAs in root.

(B) GO functional enrichment analysis for predicted target genes of miRNAs in leaf.

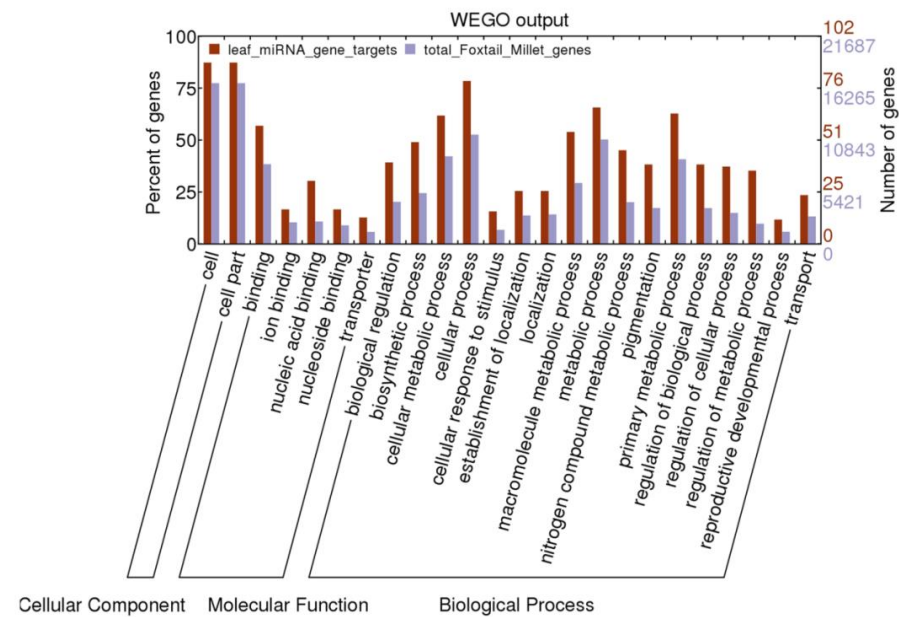
(C) GO functional enrichment analysis for predicted target genes of miRNAs in flower.

(D) GO functional enrichment analysis for predicted target genes of miRNAs in shoot.

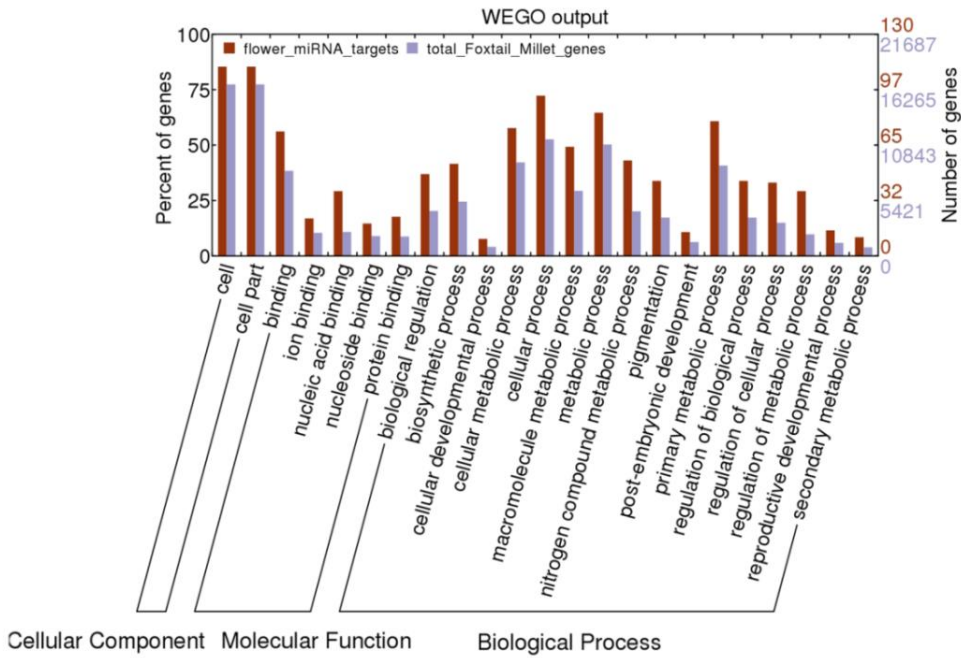
A



B



C



D

