

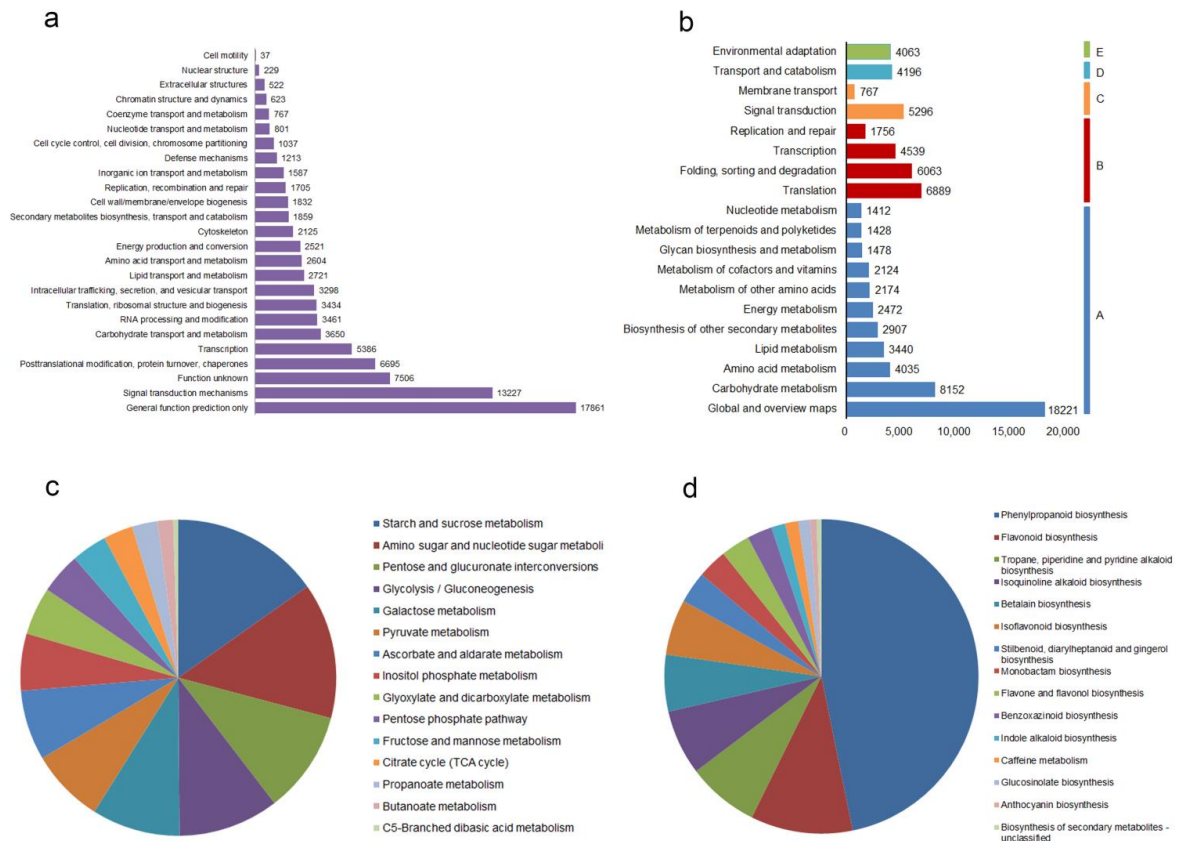


Fig. S3. Functional analysis of the assembled transcriptome. a. KOG annotation. b. Functional distribution of KEGG annotation. b-A. Metabolism; b-B. Genetic Information Processing; b-C.Environmental Information Processing; b-D. Cellular Processes; b-E. Organismal Systems. c. Sub-pathways of carbohydrate metabolism. d. Sub-pathways of biosynthesis of other secondary metabolites.