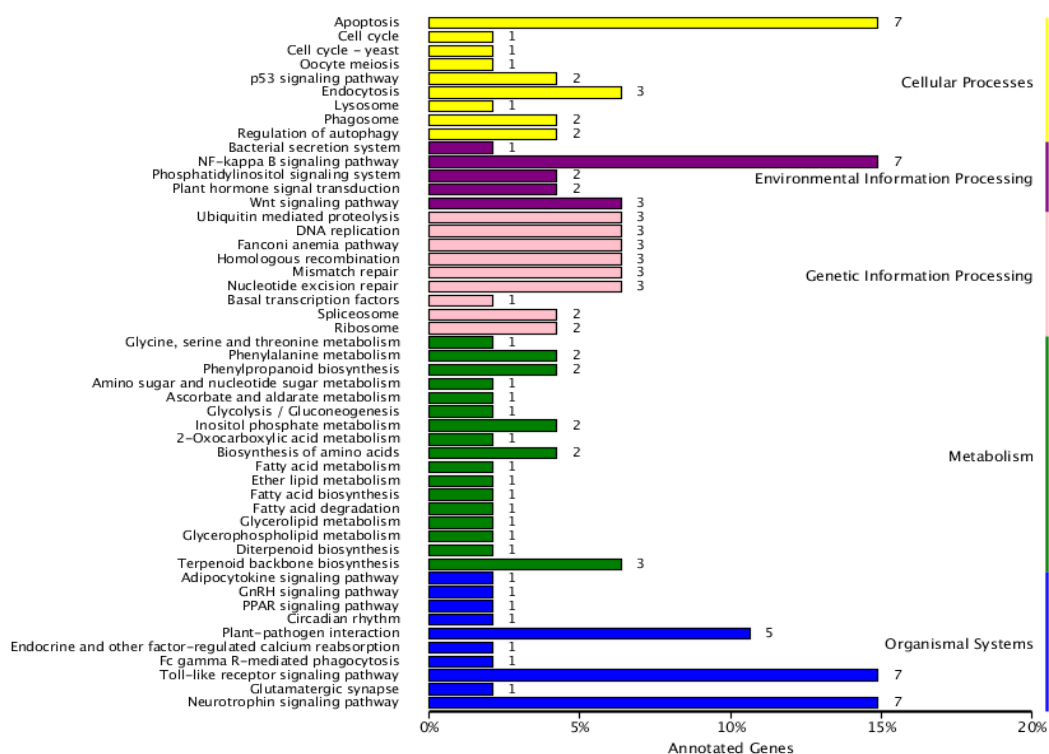
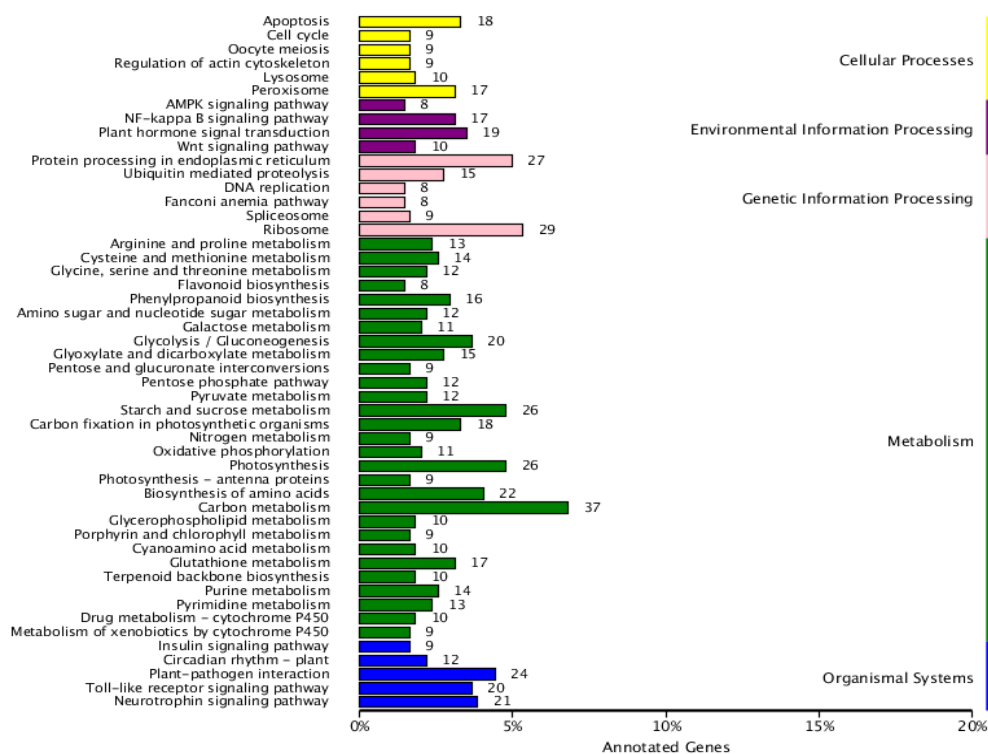
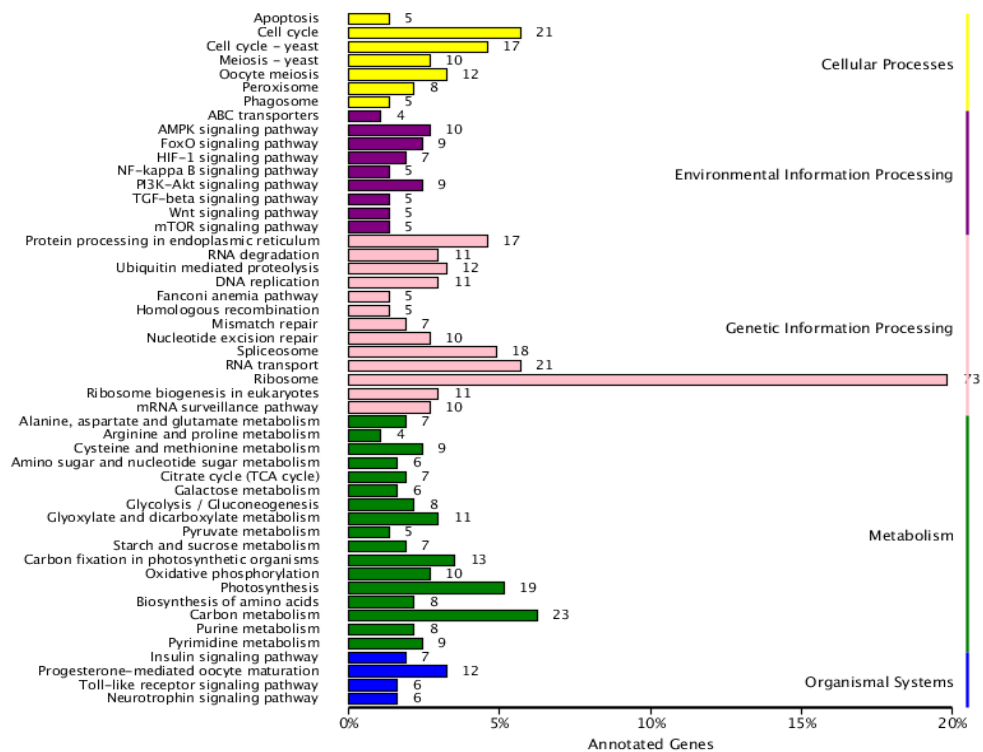




A: KEGG enrichment of DETs from XH09-7 vs ZhN03-7



B: KEGG enrichment of DETs from XH09-21 vs ZhN03-21



C: KEGG enrichment of DETs from XH09-28 vs ZhN03-28

Figure S2. KEGG classification results of differentially expressed transcripts (DETs) found in three DETs sets. All DETs were assigned to five categories: cellular process, environmental information processing, genetic information processing, metabolism and organismal systems.