Fig. S3 Directed acyclic graph (DAG) of GO enrichment analysis of candidate genes for mid-maturing resources.

Fig. 3 KEGG pathway enrichment of the candidate genes in different mature cabbages. (a, b) are the enrichment result of candidate genes located in the early-maturing (a) and mid-maturing (b) resources respectively in early-maturing versus mid-maturing analysis. (c, d) are result of candidate genes located in the mid-maturing (c) and late-maturing (d) resources respectively in mid-maturing versus late-maturing analysis. (e, f) are result of candidate genes located in the early-maturing (e) and late-maturing (f) resources respectively in early-maturing versus late-maturing analysis.

Fig. 4 GO annotation of the candidate genes in different mature cabbages. (a, b) are the annotation result of candidate genes located in the early-maturing (a) and mid-maturing (b) resources respectively in early-maturing versus mid-maturing analysis. (c, d) are result of candidate genes located in the mid-maturing (c) and late-maturing (d) resources respectively in mid-maturing versus late-maturing analysis. (e, f) are result of candidate genes located in the early-maturing (e) and late-maturing (f) resources respectively in early-maturing versus late-maturing analysis.