

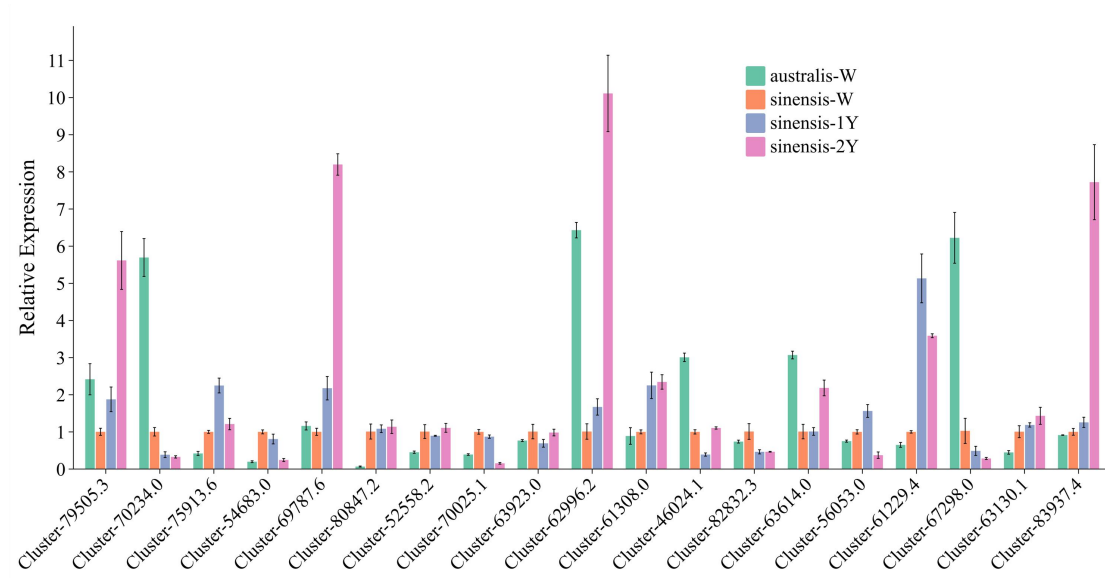


Figure S1 The relative expression values of well-annotated unigenes identified in the transcriptomic analysis. *Cluster-83937.5(acitn)* was used as an internal control. Error bars represent mean + SD of three biological replicates.