

Fig. S2

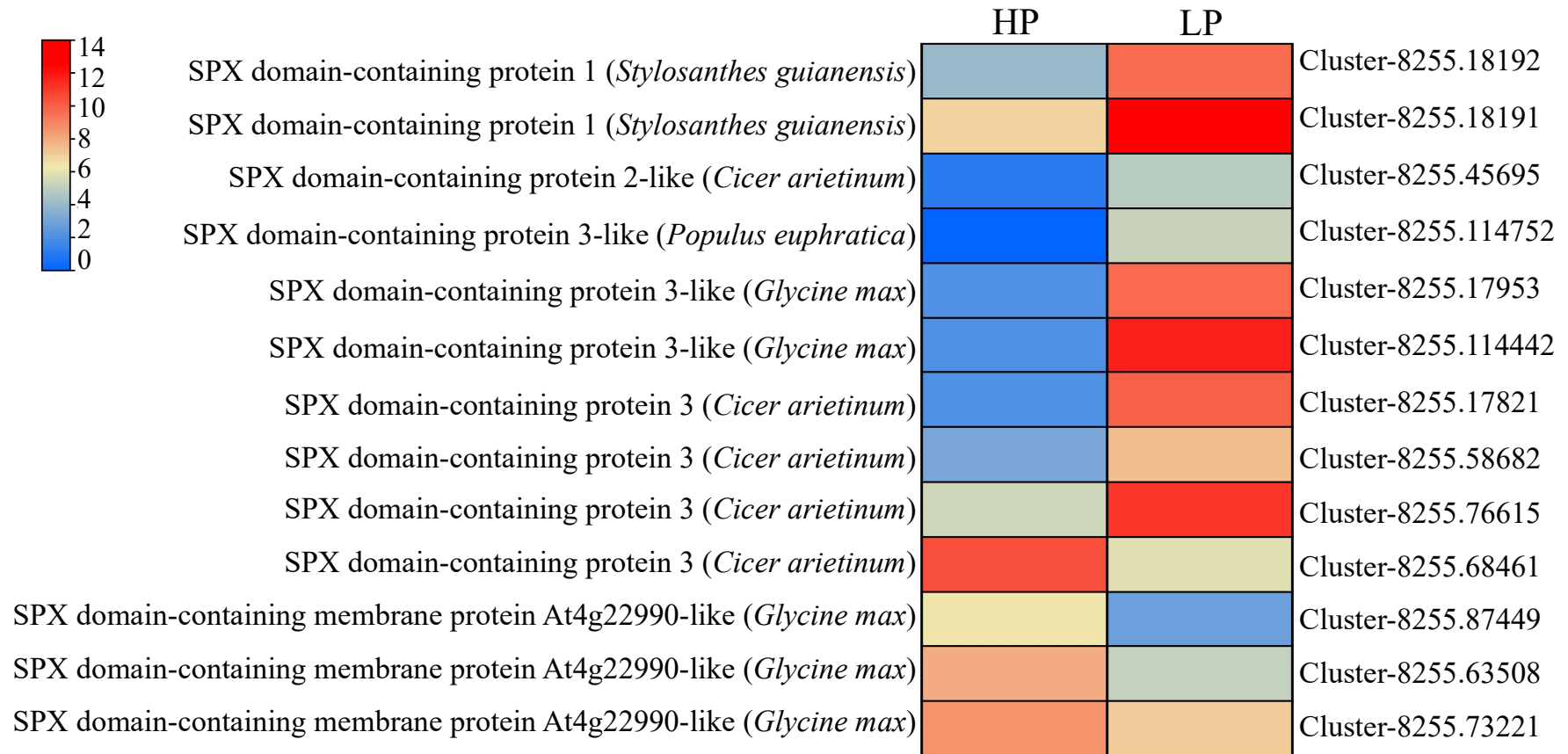


Fig. S2. Heatmap analysis of the DEGs belonging to SPX containing proteins. The transcripts of DEGs were normalized as $\log_2(\text{FPKM}+1)$. Gene IDs were showed by the legend on the right. Expression levels ranged from red to blue indicate high to low expression for genes, respectively. LP and HP represent 0 and 250 μM KH_2PO_4 supply treatments, respectively.