

Figure S1. Gene function annotations using different databases.

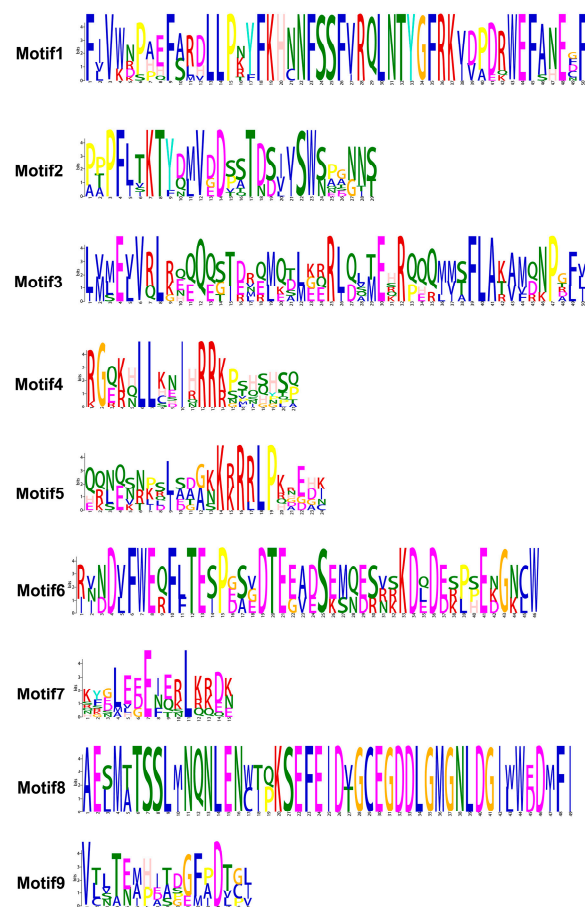


Figure S2. motif structure map of CIHsps gene family.

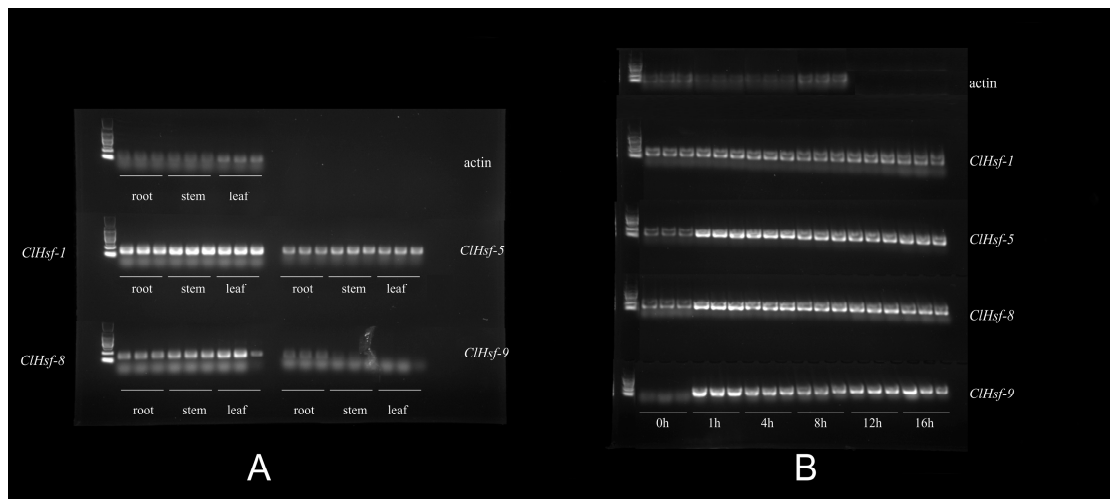


Figure S3. Semi-quantitative PCR of *CIHsf* gene family.

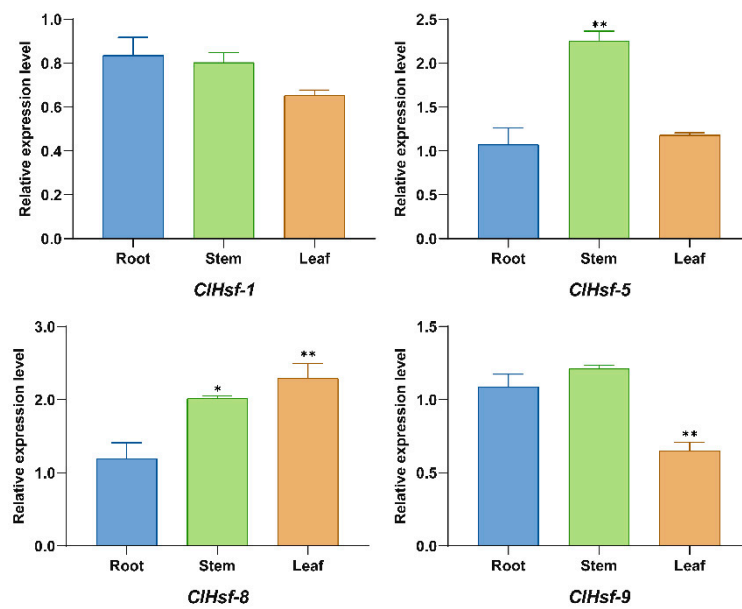


Figure S4. Expression analysis of *CIHsfs* in different tissues. Analysis of variance was used for statistical analysis. \*  $p < 0.05$ , \*\*  $p < 0.01$ , \*\*\*  $p < 0.001$ .