

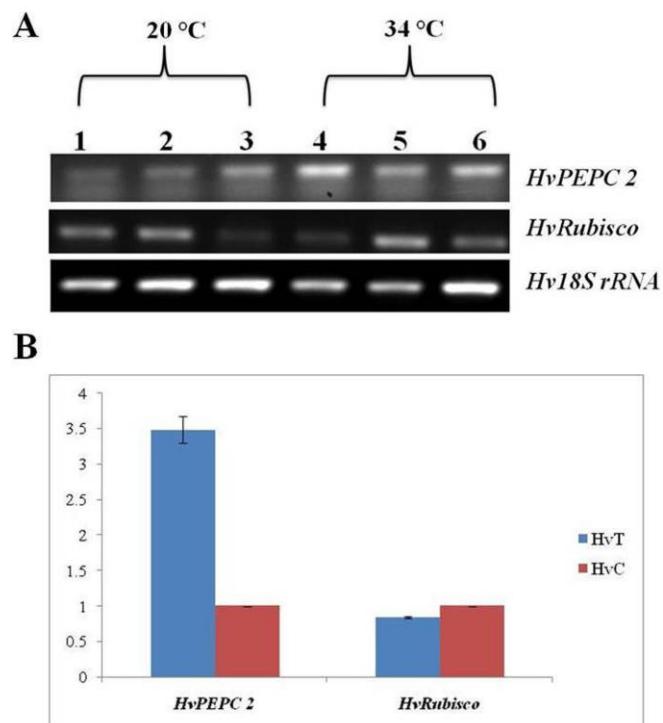
De novo transcriptome sequencing and analysis of *Hydrilla verticillata* (L.f.) Royle**Etika Goyal, Singh Amit Kumar, Ravi Shankar Singh, Ajay Kumar Mahato and Kumar Kanika***

Fig. S1: Confirmation of the transition from the C₃ to the C₄ CCM. (a) Gel electrophoresis of semi-quantitative RT-PCR using *PEPC 2* and 18s rRNA gene specific primer. Lane 1-3 D8 to D10 kept at 20°C; Lane 4-6 D8 to D10 kept at 34°C; (b) qRT-PCR expression analysis of *PEPC 2* and *Rubisco* gene expression levels.

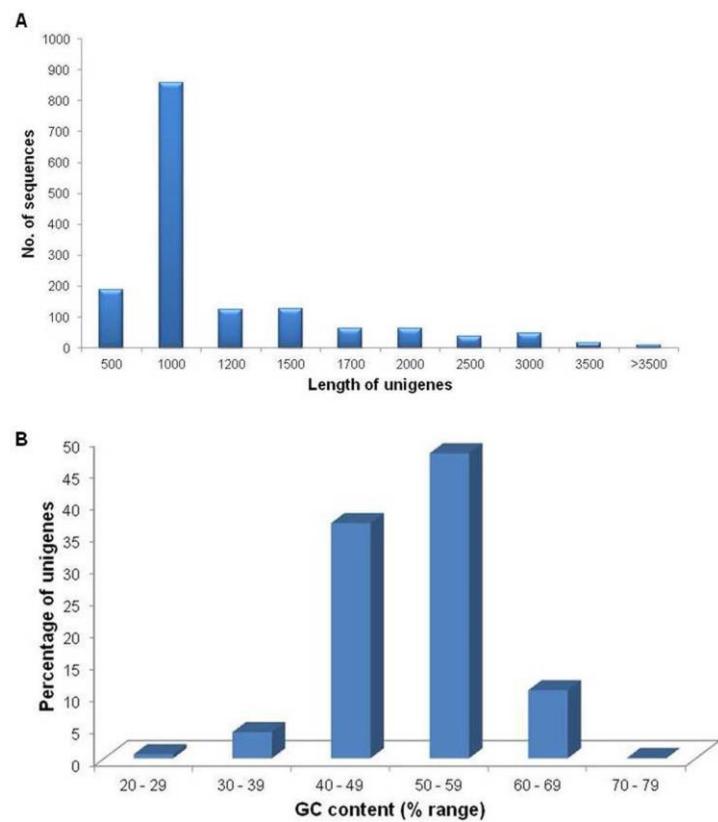


Fig. S2: Overview of *H. verticillata* transcriptome assembly. (a) Size distribution of the unigenes Y-axis: number of unigenes; X-axis: size in bp; (b) GC content of the unigenes.