



Supp. Fig. 4 qRT-PCR analysis of temporal variation in transcript abundance in all known *Eucalyptus* Cesa genes. Results were obtained for *Eucalyptus grandis* x *E. urophylla* (GU) and *E. grandis* x *E. camaldulensis* (GC) grown in the field, and under controlled light conditions. Relative expression is compared to the geometric mean of expression in three control genes. The X-axis shows Zeitgeber Time, or hours since dawn. Heavily shaded areas indicate night time and unshaded areas day time. Lightly shaded areas represent subjective night during continuous light (LL). Standard error bars represent one standard error of the mean (SEM) across three replicate measurements. The gap in the profile of CesA4 GC Field Grown corresponds to a missing value.