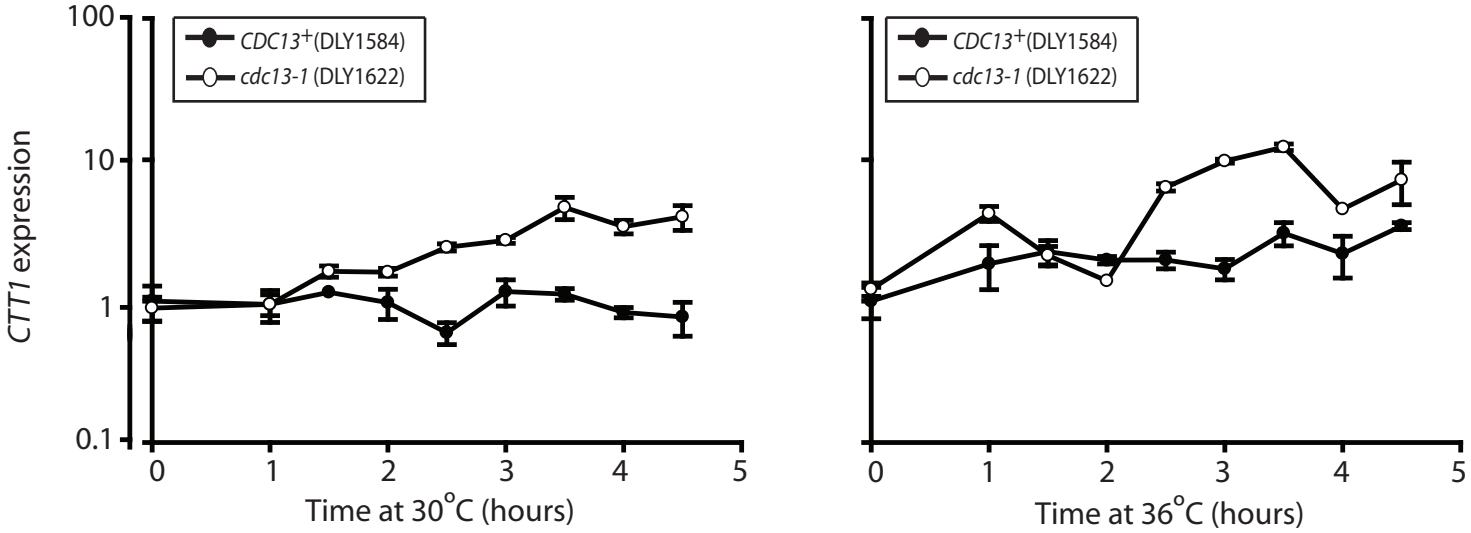
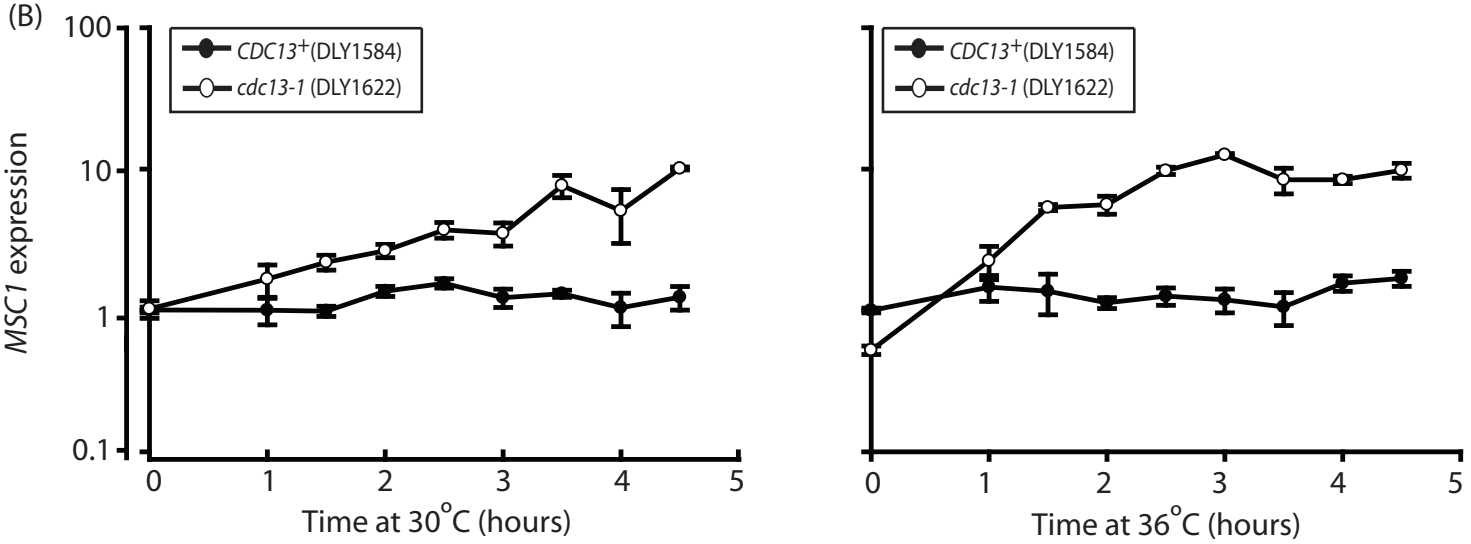
(A)



(A) *cdc13-1* (DLY1622; open circles) and *CDC13*$^{+}$ (DLY1584; filled circles) strains, grown at 23ºC, were transferred to 30ºC or 36ºC and samples taken as indicated. RNA was prepared and *CTT1* transcripts were quantified using one-step quantitative RT-PCR. Plotted values represent the means of 3 independent measurements of each sample and error bars represent the standard deviations of the means. Correction factors to normalise *CTT1* RNA concentrations of each sample were generated by calculating geometric means of three loading controls *ACT1, PAC2* and *BUD6*. A single T=0 sample from the *CDC13*$^{+}$ strain was assigned the value of 1 and all other values were corrected relative to this.

(B)



(B) Experiment carried out as in (A) but expression levels of *MSC1* were quantified.