(A) cdc13-1 (DLY1622; open circles) and CDC13+ (DLY1584; filled circles) strains, grown at 23°C, were transferred to 30°C or 36°C 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) Experiment carried out as in (A) but expression levels of MSC1 were quantified.

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