



Figure S1

Mutants of *POL30* did not increase levels of *CRE* transcripts in CRASH strains

RT-qPCR of *CRE* transcripts from *HML α 2 Δ ::*CRE* strains. Quantification was performed using a standard curve for each set of primers and normalized to *ACT1* transcript levels. Bars represent the normalized average of three technical replicates of each indicated strain: *POL30* (JRY10790), *sir4Δ* (JRY12391), *pol30-6* (JRY11139), *pol30-8* (JRY11188), and *pol30-79* (JRY11141).*