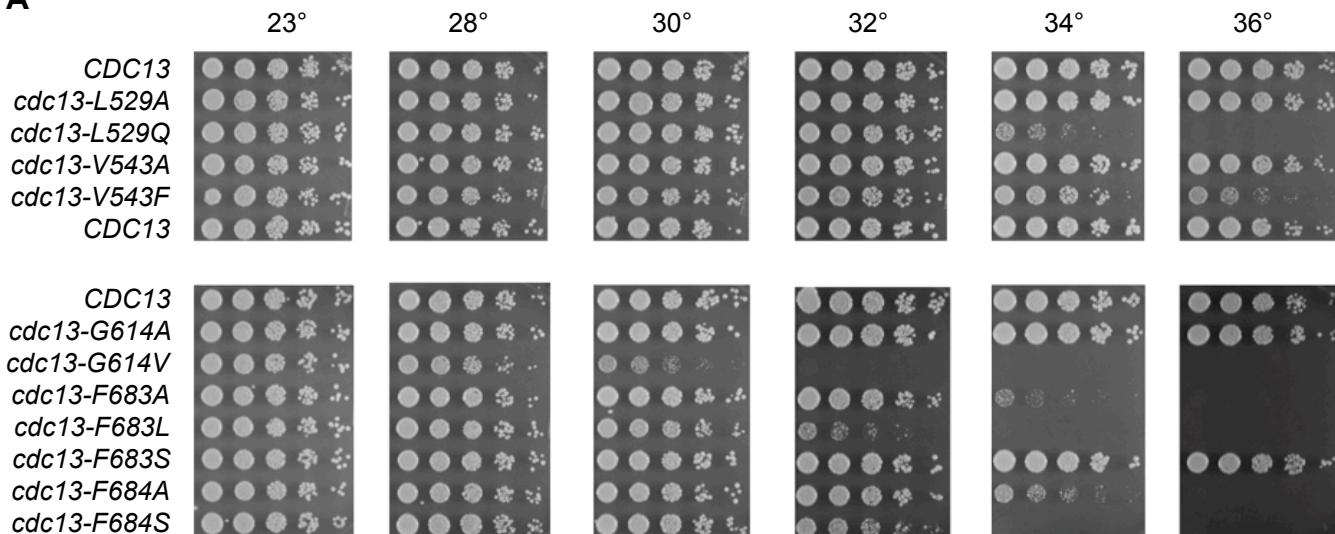
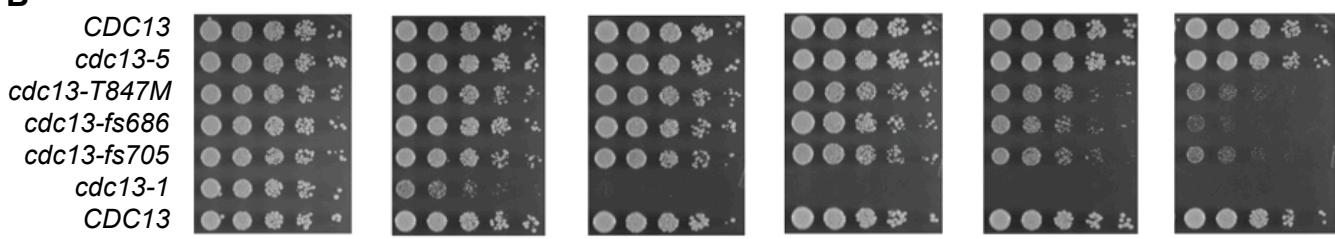


A**B****C**

Allele	Mutations present (cluster 1, cluster 2, cluster 3)
isolate 22	V543F
isolate 23	N609D
isolate 26	N609D, Y626F
isolate 19	L599P, S650P
isolate 16	M525T, I633V
isolate 30	T507A, L529Q
isolate 21	S533L, V616A
isolate 15	M463I, S611L
isolate 31	P506S, G614V
isolate 28	K618E, F683S
isolate 20	N631D, F684S
isolate 17	D535G, P651L, F683L
isolate 24	V481E, A489T, L562Q
isolate 25	T496S, E519V, I578T
isolate 27	F575L, N627S, Y680C, I686V
isolate 29	M498V, F544L, T586M, K629E
isolate 18	S531T, I563V, E608V, K629E, L693P
isolate 14	N455D, T473A, D492G, I552T, F574S

Figure S2 Identification of new *cdc13*-*ts* mutations, Part II. (A) Additional mutant alleles, as discussed in the text, were analyzed as described in Figure S1. (B) Missense mutations identified in a panel of *cdc13*-*ts* alleles, following mutagenesis of the DNA binding domain with error-prone PCR, remissense; clusters of residues discussed in the text are indicated.