

Table S1 Telomere length correction factors (telomere/rDNA)		
Tagged protein	Genetic background	Correction factor ^a
Tpz1-myc	wt	1.00 ± 0.08
	<i>poz1Δ</i>	4.72 ± 0.29
	<i>tpz1-W498R,I501R</i>	4.71 ± 0.28
	<i>poz1Δ tpz1-W498R,I501R</i>	4.86 ± 0.14
Ccq1-myc	wt	1.00 ± 0.03
	<i>poz1Δ</i>	2.63 ± 0.09
	<i>tpz1-W498R,I501R</i>	2.69 ± 0.08
	<i>poz1Δ tpz1-W498R,I501R</i>	2.63 ± 0.08
Poz1-myc	wt	1.00 ± 0.05
	<i>rap1Δ</i>	3.69 ± 0.14
	<i>tpz1-W498R,I501R</i>	3.78 ± 0.16
	<i>rap1Δ tpz1-W498R,I501R</i>	4.01 ± 0.15
Trt1-myc	wt	1.00 ± 0.02
	<i>poz1Δ</i>	3.72 ± 0.14
	<i>tpz1-W498R,I501R</i>	4.90 ± 0.14

^aMean ± standard error of the mean. Values are normalized to wild-type cells with indicated tagged proteins.