

SUPPLEMENTARY DATA

Table I. Localization of lac^{op}-tagged loci

	G1 phase				S phase			
	% in zone 1	n	p value t-test	p value χ^2	% in zone 1	n	p value t-test	p value χ^2
Tel 6R ^{nat}	59.1	110	1.3x10 ⁻⁴	6.2x10 ⁻⁸	55.6	135	2.4x10 ⁻⁴	1.4x10 ⁻⁷
Tel 14L ^{nat}	52.9	257	7.4x10 ⁻⁶	8.1x10 ⁻¹¹	58.2	153	1.3x10 ⁻⁵	6.0x10 ⁻¹⁰
Tel 7L ^{nat}	57.3	391	1.7x10 ⁻¹¹	8.1x10 ⁻²⁵	58.2	165	5.9x10 ⁻⁶	9.9x10 ⁻¹¹
Tel 8L ^{nat}	46.2	262	2.6x10 ⁻³	2.6x10 ⁻⁶	42.5	120	8.3x10⁻²	3.8x10 ⁻²
Tel 5R ^{nat}	38.6	272	0.2	5.4x10⁻²	64.8	91	2.1x10 ⁻⁵	7.7x10 ⁻³⁴
Tel 6R ^{tr}	52.4	431	1.5x10 ⁻⁸	6.4x10 ⁻¹⁷	58.5	195	6.4x10 ⁻⁷	5.6x10 ⁻¹⁴
Tel 8L ^{tr}	45.6	333	1.2x10 ⁻³	2.6x10 ⁻⁷	47.7	134	1.6x10 ⁻²	1.3x10 ⁻⁴
Tel 5R ^{tr}	50.7	276	3.5x10 ⁻⁵	4.3x10 ⁻¹⁰	56.3	112	1.8x10 ⁻⁶	1.3x10 ⁻⁴
Tel 6R ^{nat} <i>ku70Δ</i>	30	223	0.5	0.4	36.2	174	0.6	0.7
Tel 6R ^{tr} <i>ku70Δ</i>	46.7	227	3.7x10 ⁻³	7.3x10 ⁻⁴	58.8	131	3.6x10 ⁻⁵	1.2x10 ⁻⁹
Tel 8L ^{nat} <i>ku70Δ</i>	33.8	207	0.9	0.5	43.6	94	0.1	5.6x10⁻²
Tel 8L ^{tr} <i>ku70Δ</i>	53.3	214	7.1x10 ⁻⁵	1.1x10 ⁻⁸	55.2	67	1.1x10 ⁻²	3.6x10 ⁻⁴
Tel 5R ^{nat} <i>ku70Δ</i>	38.7	284	0.2	5.1x10 ⁻³	41.3	126	0.2	0.1
Tel 5R ^{tr} <i>ku70Δ</i>	45.9	231	5.8x10 ⁻³	8.1x10 ⁻⁶	57.6	118	1.8x10 ⁻⁴	6.3x10 ⁻⁸
Tel 6R ^{tr} + pAT4	57	351	3.1x10 ⁻¹⁰	1.1x10 ⁻²⁰	54.3	105	6.4x10 ⁻⁴	3.0x10 ⁻⁵
Tel 6R ^{tr} + pReb1N	56.9	239	2.3x10 ⁻⁷	9.1x10 ⁻¹⁴	50.9	53	6.6x10⁻²	1.0x10 ⁻²
Tel 6R ^{tr} + pTbf1N	46.8	314	5.7x10 ⁻⁴	2.5x10 ⁻⁶	50.6	81	2.6x10 ⁻²	1.7x10 ⁻³
Tel 6R ^{tr} <i>ku70Δ</i> + pAT4	47.4	133	2.0x10 ⁻²	2.7x10 ⁻³	59.3	86	6.4x10 ⁻⁴	2.0x10 ⁻⁶
Tel 6R ^{tr} <i>ku70Δ</i> + pReb1N	37.1	239	0.4	0.1	41.9	74	0.3	0.2
Tel 6R ^{tr} <i>ku70Δ</i> + pTbf1N	38.9	368	4.3x10 ⁻²	4.4x10 ⁻²	51.9	89	1.3x10 ⁻²	6.0x10 ⁻⁴
Tel 6R ^{tr} <i>ku70Δ</i> + pOrc2	47.5	198	4.1x10 ⁻³	7.8x10 ⁻⁴	51.9	79	1.8x10 ⁻²	1.2x10 ⁻³
Tel 6R ^{nat} <i>sir4Δ</i> + pAT4	46.2	221	5.9x10 ⁻³	8.3x10 ⁻⁵	51.1	92	1.5x10 ⁻²	2.3x10 ⁻⁴
Tel 6R ^{nat} <i>sir4Δ</i> + pReb1N	37.2	349	0.3	0.2	40.8	169	0.2	2.8x10 ⁻²
Tel 6R ^{nat} <i>sir4Δ</i> + pTbf1N	43	207	4.3x10 ⁻²	7.7x10 ⁻³	50.6	87	2.1x10 ⁻²	3.0x10 ⁻³
Tel 6R ^{tr} + pVP16	37.6	234	0.3	0.2	37.7	85	0.6	0.3
Tel 8L ^{nat} + pVP16	46.6	148	2.0x10 ⁻²	6.4x10 ⁻⁴	54.4	57	2.4x10 ⁻²	3.4x10 ⁻³
Tel 6R ^{nat} <i>tell1Δ</i> + pAT4	51.6	184	3.9x10 ⁻⁴	5.8x10 ⁻⁷	49.3	69	5.7x10⁻²	1.7x10 ⁻³
Tel 6R ^{nat} <i>tell1Δ</i> + pReb1N	36.9	306	0.4	0.4	43.0	121	0.1	3.8x10 ⁻²
Tel 6R ^{nat} <i>tell1Δ</i> + pTbf1N	43.9	312	6.7x10 ⁻³	2.9x10 ⁻⁴	47.0	83	5.5x10⁻²	2.1x10 ⁻³
Tel 6R ^{nat} <i>tell1Δ</i> + pVP16	39.3	145	0.3	0.1	45.5	55	0.2	5.0x10 ⁻³
Tel 6R ^{tr} <i>tell1Δ</i> + pAT4	51.6	184	1.9x10 ⁻³	2.2x10 ⁻⁵	49.3	69	0.1	8.0x10⁻²
Tel 6R ^{tr} <i>tell1Δ</i> + pReb1N	36.9	306	0.9	0.4	43.0	121	0.5	6.3x10⁻²
Tel 6R ^{tr} <i>tell1Δ</i> + pTbf1N	43.9	312	0.5	8.1x10⁻²	46.0	111	0.4	0.2
Tel 6R ^{tr} <i>tell1Δ</i> + pVP16	39.3	145	0.2	7.9x10 ⁻³	45.5	55	0.3	0.2

n represents the total number of cells counted for a specific condition, the p value is obtained with either a t-test comparing the proportion of total spots in zone 1 to the 33% theoretical random proportion, or with a Chi-squared test comparing telomeres in zones 1, 2 and 3 to a theoretical random distribution with an equal n. p values > 0.05 are in bold and indicate no significant difference from random distributions.