

Table S18. Base substitutions within complex mutations.

WT base (in unresected strand)	<i>can1</i> no-DSB; UV (45) (noncoding)	<i>can1</i> DSB- <i>cen</i> ; no UV	<i>can1</i> DSB- <i>cen</i> ; UV (20)	<i>can1</i> DSB- <i>cen</i> ; UV (45)	<i>can1</i> (<i>can1 ura3</i>) DSB- <i>cen</i> ; UV (45)	<i>ura3</i> (<i>can1 ura3</i>) DSB- <i>cen</i> ; UV (45)	<i>can1</i> DSB- <i>tel</i> ; UV (45)	subtel. <i>LYS2</i> 23°C (no arrest) UV (45) (coding)	subtel. <i>LYS2</i> 37°C (arrest) UV (45)
Pu	7					1			1
Py	12	2	10	6	3	6	2		6
Total complex mutations¹	11	1	6	3	2	3	1	0	5

¹ Includes all complex mutations regardless of the number and types of changes within a complex mutation.

Also see footnotes to Table S16.