

Table S2. Loci used in the phylogeny of CTG yeast species

Annotation in <i>S. cerevisiae</i>	Description
MS456 (NP_009761)	Mcm7, Mcm2-7 helicase subunit
FG533 (NP_015239)	Elp3, Elongator subunit
MS378 (NP_012056)	Kog1, Ubiquitin-binding TORC1 subunit
FG570 (NP_010066)	Gdh2, Glutamate dehydrogenase
FG595 (NP_013826)	Ilv2, Acetolactate synthase catalytic subunit
FG848 (NP_012712)	Uba1, Ubiquitin activating E1 protein
FG692 (NP_014247)	Pol1, catalytic subunit DNA polymerase alpha
FG610 (NP_012526)	Cct8, Chaperonin Cct ring complex subunit
MS407(NP_010476)	Rvb1, RUVB family ATP-dependent DNA helicase
MS431(NP_010016)	Msh3, Mismatch repair protein