

Table S7. Origin and number of TCsat12, TCsat13, and TCsat14 monomers used in phylogenetic analyses.

	TCsat12	TCsat13	TCsat14
<i>Cloned PCR-dimers</i> (containing one full-size monomer)	13	24	12
<i>Cloned PCR-trimers</i> (containing two full-size monomers)	9	5	17
Extracted PCR monomers	31	34	46
Monomers from assembled contigs	4	-	36
Monomers from unplaced scaffolds	7	19	1
Monomers from unplaced singletons	71	31	-
Total number of analyzed monomers	113	84	83