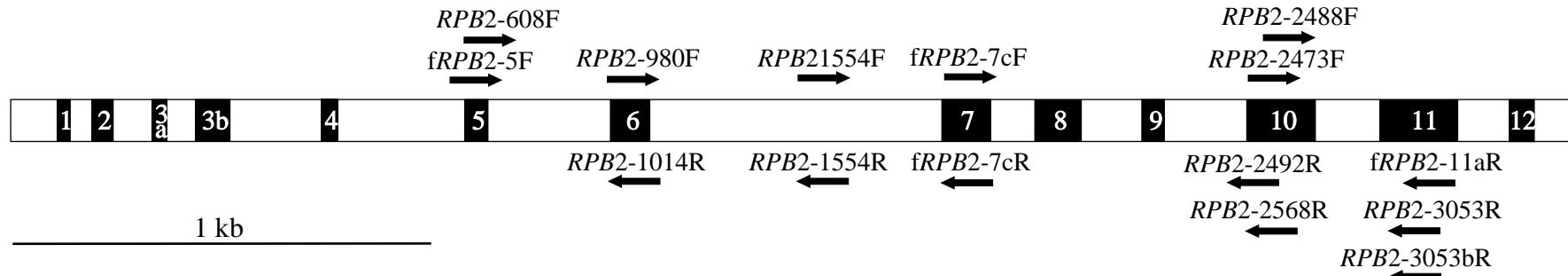


## *RPB2*



<u>Primer name</u>	<u>Sequence</u>	<u>Position</u>	<u>Reference</u>
f <i>RPB2</i> -5F	5'-GAYGAYMGWGATCAYTTYGG-3'	1186-1205	Liu et al. (1999)
f <i>RPB2</i> -7cF	5'-ATGGGYAARCAAGCYATGGG-3'	2317-2336	Liu et al. (1999)
f <i>RPB2</i> -7cR	5'-CCCATRGCTTGYTRCCCAT-3'	2336-2317	Liu et al. (1999)
f <i>RPB2</i> -11aR	5'-GCRTGGATCTTRTCRTCSACC-3'	3314-3294	Liu et al. (1999)
R <i>RPB2</i> -608F	5'-GAYCAYTTYGGIAARAA-3'	1195-1211	Reeb et al. (2004)
R <i>RPB2</i> -980F	5'-TGYCCIGCIGARACICCHGARGG-3'	1567-1589	Reeb et al. (2004)
R <i>RPB2</i> -1014R	5'-CCRCAIGCYTGICCYTCDDGG-3'	1601-1582	Reeb et al. (2004)
R <i>RPB2</i> -1554F	5'-GAYGCIGARGARGARGA-3'	2080-2096	Reeb et al. (2004)
R <i>RPB2</i> -1554R	5'-TCYTCYTCYTCIGCRT-3'	2096-2080	Reeb et al. (2004)
R <i>RPB2</i> -2473F	5'-CCICARATYGGIGAYAARTT-3'	2920-2939	Reeb et al. (2004)
R <i>RPB2</i> -2488F	5'-AARTTYGCITCYCGICAYGG-3'	2935-2954	Reeb et al. (2004)
R <i>RPB2</i> -2492R	5'-AAYTTRTCICC RATYTG-3'	2939-2923	Reeb et al. (2004)
R <i>RPB2</i> -2568R	5'-TAIGTRAYACCRATIGTWCC-3'	2981-2962	Reeb et al. (2004)
R <i>RPB2</i> -3053R	5'-TGRATYTTRTCRTCSACCATRTG-3'	3311-3289	Reeb et al. (2004)
R <i>RPB2</i> -3053bR	5'-TGRATYTTRTCRTCSACCAT-3'	3311-3292	Reeb et al. (2004)

Supplementary material 2 - Reeb et al. (2004) Mol. Phylogenetic Evol. 32:1036-1058

Representation of the *RPB2* gene according to Liu et al. (1999). The black numbered boxes represent the 12 amino acid regions that are conserved throughout eukaryotes. Positions of the *RPB2* primers used for symmetric PCR amplifications and cycle sequencing reactions are represented by black arrows. Primer names in the table are followed by their degenerate oligonucleotide sequences, their positions relative to the *RPB2* gene sequence of *Saccharomyces cerevisiae* (GenBank Accession No. M15693), and their respective sources.