

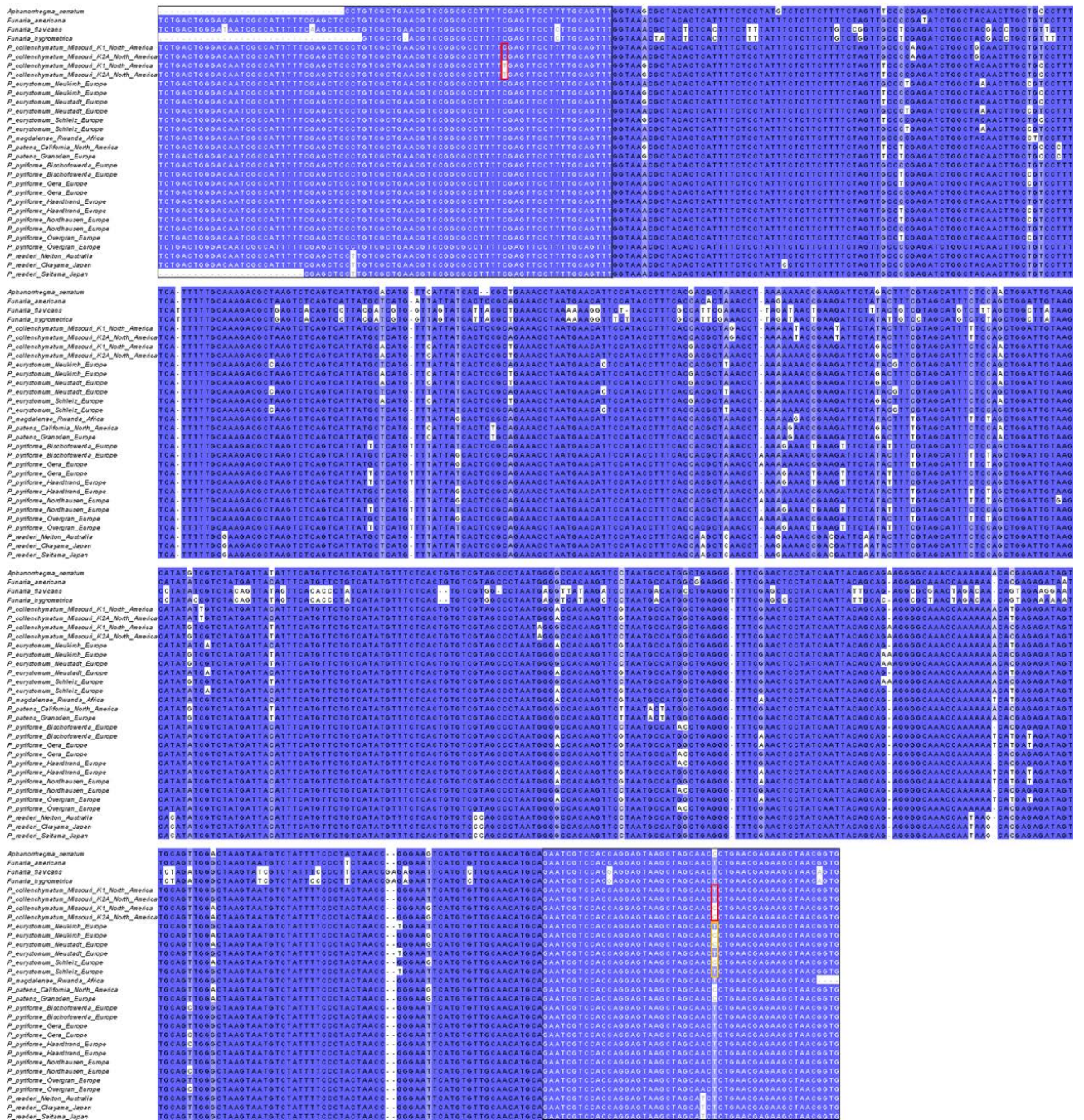


Figure S1 – Alignment of genomic *BRK1* sequences

Multiple sequence alignment of amplified and clonal genomic sequences of *BRK1* from different Funariaceae. The exon region is shown in white letters, the intron region in black letters. The species names are sorted alphabetically. Accessions with only one locus of *BRK1* are represented by one sequence of directly sequenced PCR product, whereas two representative clonal sequences are shown for each accession with two loci of *BRK1* (*P.*

collenchamytum, *P. eurystomum*, and *P. pyriforme*). Polymorphisms in the exon regions are depicted in red (*P. collenchymatum*) and orange (*P. eurystomum*) boxes.