

Sanger sequencing results of *CG10212_r*, *CG33957_r* and *CG4799_CG11924_r*.
 The information about locations and primers can be found in Supplemental Table 6.
 The ‘p_LTR’ sequence is artificially constructed on the basis of the parental gene and LTR retrotransposon sequences encoded in the iso-1 genome.

Sequence alignment

The alignment is color-coded as follows:

5’ and 3’ LTR retrotransposon repeat zone

Coding region of LTR retrotransposon *BLASTOPIA*

Microsimilarity shared by LTR retrotransposon and retrocopy

Retrocopy

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