

A *GPD1*

M. amblycephala genome
Chromosome 10

Sequence by Sanger method

***C. alburnus* genome**
Contig197

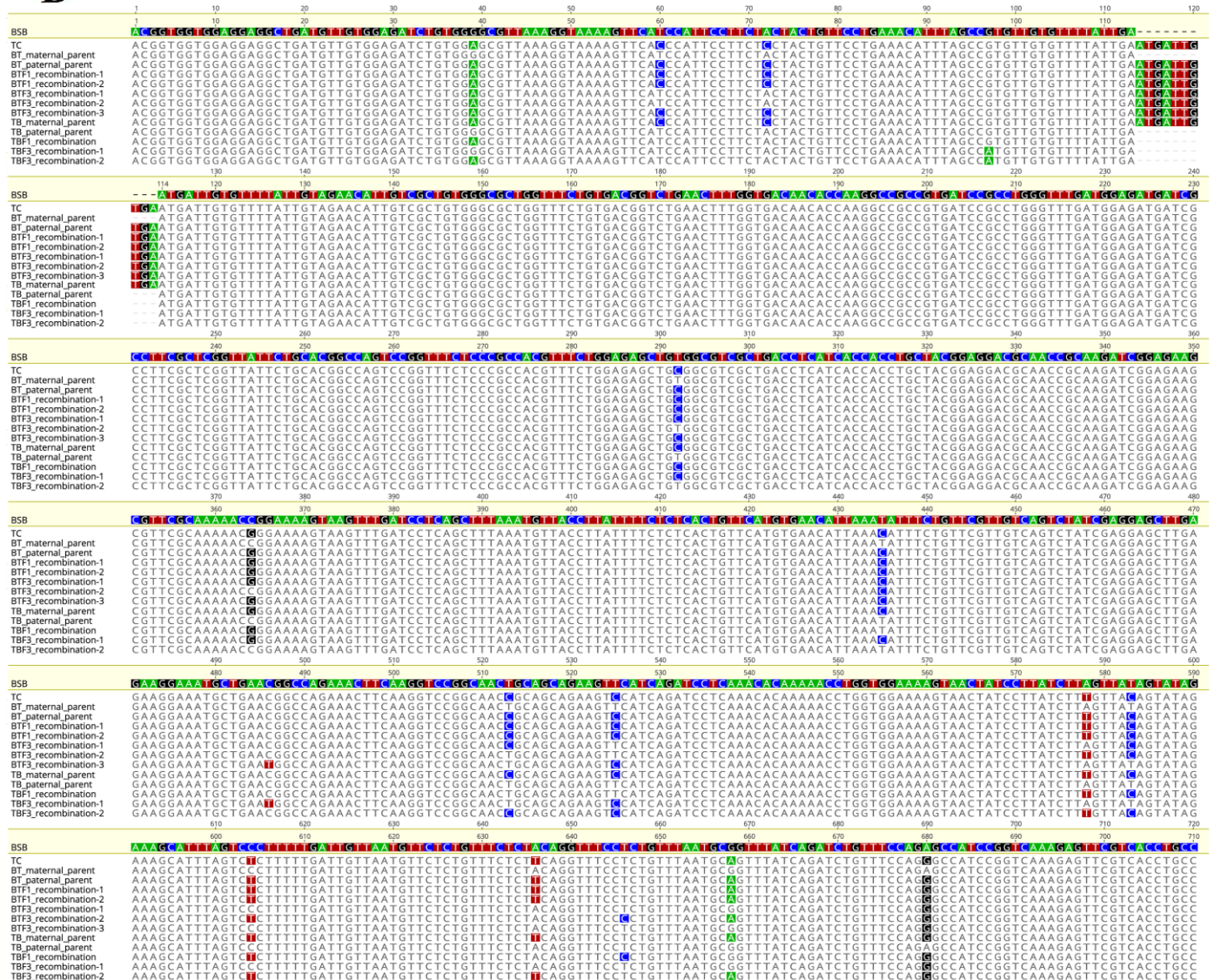
6,726,407

6,727,118

181,362

182,083

B



Supplemental Fig. S10. Diagram and alignment of *GPDI* (DNA) from BSB, TC, and their reciprocal cross hybrids (BT and TB) in the F₁ and F₃ generation. (A) Schematic map of relationships between the reference genome and Sanger sequences. (B) Sequence alignment for *GPDI*.