

A EPHX2

M. amblycephala genome
Chromosome 17

Sequence by Sanger method

C. alburnus genome
Chromosome 17

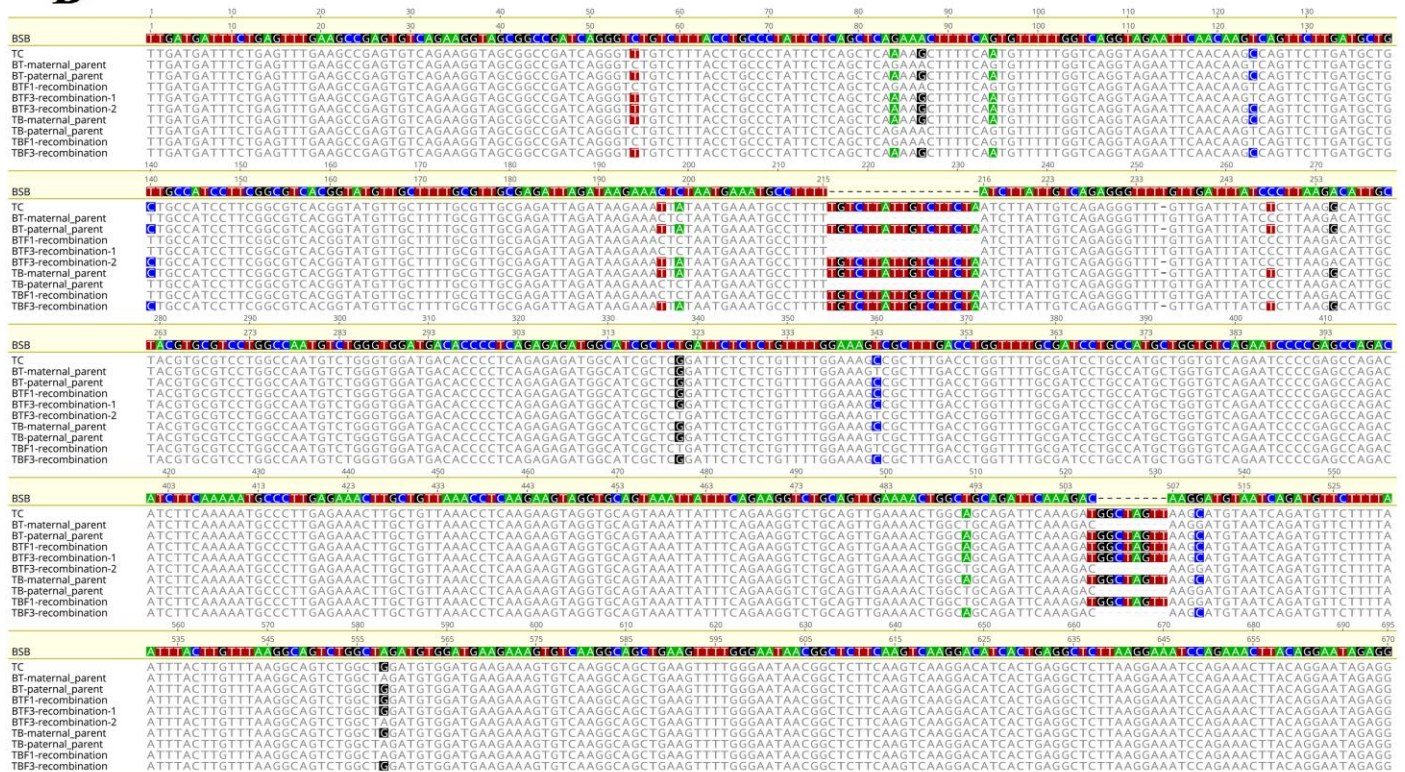
7,411,384

7,412,052

1,088,979

1,089,671

B



Supplemental Fig. S22. Diagram and alignment of *EPHX2* (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 *EPHX2*.