Figure S1: Sequence alignment of the 367 bp-left-end (A) and 230 bp-right-end (B) silver carp Thm1 with blunt snout bream Tma2. The nested primers for the 5’ or 3’ flanking sequences are marked in gray. Dashed lines represent missing nucleotides and dots mean similar nucleotides. The double underlined sequences are the end sequences of Thm1 DNA at target integration sites in the genome of the blunt snout bream as shown in Table 3.