Fig. S7. Information on the polymorphic site YM23 - Pos 4161

(a) Sequence alignment (HT1,2,3,4,5,6; HTPrz1; HTPrz2; Reference from BAC clone E) with primer positions underlined. Positions identical to the sequence in the first raw are represented with a hyphen. The mutation leading to HTPrz2 is marked. (b) PCR products amplified from male (m) and female (f) genomic DNA and a no-template-control (-). (c) Capillary sequence traces of the confirmed mutations in HT1,2,3,4,5,6 and HTPrz1 and HTPrz2 respectively.

