

5' -TTTTTTTTAAAGTAGCCTTTCTTATGTTCTTTCTTTGTTATATATTT
CTGGCCCTGCTGCTCCGGCCTTATTATTTATAAATGTGAGTACACTATTGCTCTTCAGA
TACCAGAACAGGGCACCAAATCCCATTATAGATGGTTGTGAGCCACCATGTGGTTGCTGGCGAA
AGGGGGATGTGCTGCAAGGCAGTAAGTTGGTAACGCCAGGGTTTCCCAGTCACGACGTTGT
AAAACGACGGCCAGTGAATTGAGCTCGGTACCCATCCTGCCTTAATGCCTTGATGCATGTA
TACAAGCTAACACAGGCTTCACTTCTGCCAACTTACAAGGCCTTCTAAGTAAACAGTACAT
GAACCTTACCCGTTGCTCGCAACGGCCTGGTCTGTGCCAAGTGTGCTGACGCAACCCCC
ACTGGCTGGGCTTGGCATAGGCCATCAGCGCATGCGTGGAACCTTGTCGGC-3'

Figure S6. Sequences at the junction of HBV (red) and mouse genomic (green) DNA in Tg05 wildtype HBV transgenic mice. The vector sequence is in black. The data is derived from sequencing the PCR product shown in Figure 6G. The DNA sequence in green is identical to 78,654,987-78,654,805 of mouse chromosome 11 and is in 11qB5 region (UCSC Genome Browser).