

5' -CTGCTGGGTGACCTGGCTGCCCTGCCCGCCGATCTGAGAACCCGTAAGCGGGATCTTGACC
GCGGGGAATGGTATCTGGAGCCCGCAAACCTGCCGCTCAGCGGGTATACATTTAAACCCTAACA
AAACAAAAGATGGGGTTATTCCTAAACTTCATGGGTACATAATTGGAAGTTGGGGAACGTT
GCCACAGGATCATATTGTACAAAAGATCAAACACTGTTTTAGAAAACCTCCTGTTAACAGGCCT
ATTGATTGGAAAGTATGTCAAAGAATTGTGGGTCTTTTGGGCTTTGCTGCTCCATTTACACAAT
GTGGATATCCTGCCTTAATGCCTTTGTATGCATGTATAACAAGCTAAACAGGCTTTCACCTTCTC
GCCAACTTACAAGGCCTTTCTAAGTAAACAGTACATGAACCTTTACCCCGTTGCTCGGCAACGG
CCTGGTCTGTGCCAAGTGTGGCTGACGCAACCCCACTGGCTGGGGCTTGCCATAGGCCATC
AGCGCATGCGTGGAACCTTTGTGGCTCCTCTGCCGATCCATACTGCGGAACTCCTAGCCGTTGT
TTTTGCT-3'

Figure S4. Sequences at the junction of HBV (red) and mouse genomic (green) DNA in Mutant 1 Line-7 mice. The data is derived from sequencing the PCR product shown in Figure 6A. The DNA sequence in green is identical to 106,650,370-106,650,471 of mouse chromosome 11, which is located in 11qE1 region (UCSC Genome Browser).