

5' -ACCCATTAAAAACGGGGCTCAGAACTGAACAAAGAATTCTCACCTGAGGAATACCGAAT
GGCAGAGAAGCACCTGAAAAATGTTCAACATCCTTAATCATCAGGGAAATGCAAATCAAAACA
ACCCTGAGATTCTACCTCACACCAGTCAGAATGGCTAAGGGGGATGTGCTGCAAGGCGATTAAG
TTGGGTAACGCCAGGGTTTTCCAGTCACGACGTTGTAAAACGACGGCCAGTGCCAAGCTTGCA
TGCCTGCAGGTCGAATCCTGCCTTAATGCCTTTGTATGCATGTATACAAGCTAAACAGGCTTTC
ACTTTCTCGCCAACTTACAAGGCCTTCTAAGTAAACAGTACATGAACCTTTACCCCGTTGCTC
GGCAACGGCCTGGTCTGTGCCAAGTGTGCTGACGCAACCCCCACTGGCTGGGGCTTGCCAT
AGGCCATCAGCGCATGCGTGGAACCTTTGTGGCTCCTCTGCCGATCCATAC-3'

Figure S5. Sequences at the junction of HBV (red) and mouse genomic (green) DNA in Mutant 1 Line-4 mice. The vector sequence is in black. The data is derived from sequencing the PCR product shown in Figure 6D. The DNA sequence in green shares 99.4% identity with 145980892-145980732 of mouse chromosome 1, part of a LINE1 retrotransposon in the 1qF region (UCSC Genome Browser).