DRUG-RESISTANT MUTATION 1

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Input

Consensus

Consensus

Input

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Input

Consensus

Consensus

Input

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Input

Consensus

Consensus

Input

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Sequence Alignment

Alignment between the input query sequence and the genotype D consensus reference sequence. Amino acid and nucleotide sequence numbers refer to HBV polymerase numbering.

Figure S1. Drug-resistant mutation observed in a case of HBV-related HCC from South India. The input sequence (yellow) bears an amino acid I in place of the expected L, as observed in the reference sequence (green) at codon 80 (blue). The mutation L80I is a lamivudine mutation and a compensatory mutation for M204V/I.