

Electronic Supplementary Material

A novel phenotypic assay of hepatitis B virus polymerase with extensive site-specific mutagenesis

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Table S1. Primer sequences

Primer	5'–3'
lenti6744F	AGGTTTCCCGACTGGAAAGC
Rpol204	TTGGCCCCAATACCACATCATCNNTA TAACTGAAAGCCAAACAGTGGG GGA
F260	TAGACTCGTGGTGGACTTCTCT
RlentiPCR	GAATCGAGACCGAGG AGAGGGT
F1927	GGAGCTACTGTGGAGTTACTCT
R2051	TTGCCTGAGTGCAAGTATGGTGA

Table S2. Distribution of mutations among the 30 plasmids sequenced

Nucleotides (ANN)	Amino acid	Number of clones
AUG	M204 WT	3
AUA	M204I	1
AUU	M204I	1
AUC	M204I	2
ACA	M204T	5
ACU	M204T	1
ACC	M204T	2
ACG	M204T	0
AAA	M204K	3
AAG	M204K	1
AAU	M204N	1
AAC	M204N	2
AGA	M204R	4
AGG	M204R	1
AGC	M204S	2
AGU	M204S	1
Total		30