



Supplementary Figure 1. Phylogenetic analysis of the region between nucleotides 1255 and 1611, extracted from 102 full-length HBV genome patterns, used for genotyping.

The representation of the phylogenetic analysis shows that the different patterns clustered according to their HBV genotype, which coincided with that previously determined using the full HBV genome sequence. Each color box highlights the sequences that correspond to a specific genotype (GenBank accession number reported on the right).

Supplementary table 1. Reference sequences used for genotyping

NCBI GenBank accession numbers of the 105 reference sequences used in HBV genotyping:

HBV genotype (N sequences)	GenBank accession numbers
A (13)	AY233278, AB241115, GQ477501, AF090841, AF090839, AB194952, AB194951, AM180623, AY934764, FJ692609, FJ692613, GQ331047, GQ331048
B (20)	AB073858, AB362933, D00329, AY596111, AP011084, GQ924653, M54923, AP011085, AB073835, AB115551, AB219427, AP011086, AB287316, DQ463787, EF473977, AP011091, AP011093, AP011094, GQ358148, GQ358152
C (23)	AB112066, X52939, AF533983, AB033553, X75656, X75665, AB048705, AB048704, AP011099, AB241109, AP011102, AP011103, EU670263, AP011107, AP011104, AP011108, AB540583, AB554019, AB554025, AB644281, AB644284, AB644286, AB644287
D (17)	AB555496, GU456647, AB104712, AF280817, AB210820, X97848, Z35716, EU939680, AY233291, AJ132335, GQ922005, AB048701, DQ315779, AB033558, AB493846, FJ904405, AM494716
E (8)	X75664, X75657, FJ349237, AM494694, HM363569, FJ349226, DQ060828, JQ000008
F (10)	AY090459, HQ378247, HE981184, AY090455, AY311369, X69798, X75663, AB036911, AB166850, DQ823090
G (6)	EF464098, HE981172, HE981176, AP007264, GU565217, AB064311
H (5)	AP007261, AB275308, AY090460, AY090454, AB516393