

**Supplementary Table S2: HBV genotype dependent sequences of ‘Region 1’ and ‘Region 2’.**

Hepatitis B Virus Genotype	Sequence of the viral genome regions where stemloop sequences of miRNAs are mostly aligned	
	Region 1	Region 2
<b>A</b>	GATCCTTCGCGGGACGTCCTTTGTTTACGTCCCGTCGGCGCTGAATCCCGC GGACGACCCCTCTCGGGGCC	TGTTTAAGGACTGGGAG_GAGCTGGGGGAGGAGATTAGGTTAAT GATCTTTGTATTAGTAAATTG
<b>B</b>	GATCCTGCGCGGGACGTCCTTTGTCTACGTCCCGTCGGCGCTGAATCCCG CGGACGACCCCTCCCGGGGCC	TGTTTACTGAGTGGGAG_GAGTTGGGGGAGGAGATCAGGTAA AGGTCTTTGTACTAGTAAATTG
<b>C</b>	GATCCTGCGCGGGACGTCCTTTGTCTACGTCCCGTCGGCGCTGAATCCCG CGGACGACCCGTCTCGGGGCC	TGTTTAAGGACTGGGAG_GAGTTAGGGGAGGAGACTAGGTAA TGATCTTTGTACTAGTAAATTG
<b>D</b>	GATCCTGCGCGGGACGTCCTTTGTTTACGTCCCGTCGGCGCTGAATCCCGC GGACGACCCCTCTCGGGGCC	TGTTTAAGGACTGGGAG_GAGTTGGGGGAGGAGACTAGATTAAT GATCTTTGTACTAGTAAATTG
<b>E</b>	GATCCTGCGAGGGACGTCCTTTGTCTACGTCCCGTCAGCGCTGAATCCTG CGGACGATCCGTCTCGGGGTC	TGTTTAAAGACTGGGAG_GAGTTGGGGGAGGAGATTAGATTAA AGATCTTTGTACTAGTAAATTG
<b>F</b>	GATCCTGCGCGGGACGTCCTTTGTTTACGTCCCGTCGGCGCTGAATCCCGC GGACGACCCCTCCCGGGGTC	TATTTAAGGACTGGGAG_GAGCTGGGGGAGGAGATCAGGTAA AGGTCTTTGTACTAGTAAATTG
<b>G</b>	GATCCTTCGCGGGACGTCCTTTGTTTACGTCCCGTCAGCGCTGAATCCAGC GGACGACCCCTCCCGGGGCC	TTTTTGCTGAGTGGGAA_GAATTAGGCAATGAGTCCAGGTAA GACCTTTGTATTAGTAAATTG
<b>H</b>	GATCCTGCGCGGGACGTCCTTTGTCTACGTCCCGTCGGCGCTGAATCCTGC GGACGACCCCTCTCGTGGTC	TATTTAAGGACTGGGAG_GAGTCGGGGGAGGAGTTGAGGTAA AGGTCTTTGTACTAGTAAATTG