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# Recommended Protocol for HBV genotyping and for antiviral resistance analysis.

This amplification protocol is based on a nested PCR. The final amplicon is approximately 1kb and will cover the entire HBsAg region as well as domains A to E of the overlapping polymerase region.

All primer sequences shown are 5'-3'.

Outer sense:

Outer antisense:

HBV Z - AGC CCT CAG GCT CAG GGC ATA

HBV 3 - CGT TGC CKD GCA ACS GGG TAA AG

Inner sense:

HBV P - TCA TCC TCA GGC CAT GCA GT

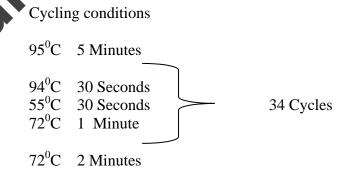
HBV M- GAC ACA CTT TCC AAT CAA TNG

### **First Round PCR**

PCR Master mix per sample: (All reagents are from Invitrogen

2.5µl	10XPCR Buffer
0.75µl	50mM Mg Cl
0.5µl	10mM dNTPS
0.1µl	Taq Polymerase
0.5µl	HBV Z (20pmol/µl)
0.5 μl	H <b>ŠΨ3</b> (20pmol/μl)
15.15µl	dH <sub>2</sub> O
0.5μl 0.5 μl	HBV 3 (20pmol/μl) HBV 3 (20pmol/μl)

Add 5µl viral DNA to 20µl PCR Master Mix.



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#### Second Round PCR

PCR Master mix per sample:

5µl	10XPCR Buffer
1.5µl	50mM Mg Cl <sub>2</sub>
1µl	10mM dNTPS
0.2μ1	Taq Polymerase
1µl	HBV P (20pmol/μl)
1µl	HBV M (20pmol/μl)
39.3µl	$dH_2O$

8.3 Add 1µl first round product to 49µl PCR Master Mix.

Cycling conditions

7 Minute

Per John 95°C 5 Minutes  $94^{0}$ C 30 Seconds 34 Cycles  $50^{0}$ C 30 Seconds  $72^{0}C$ 1 Minute  $72^{0}$ C

## **Sequencing PCR**

Sequencing reactions shoul et up in accordance with the instructions provided by the manufacturer.

and round amplicon is analysed using each of the following NB – Each sample ndividual reactions:

TCC TCA GGC CAT GCA GT

SAC ACA CTT TCC AAT CAA TNG G

IBV N- ACTGAGCCAGGAGAAACGGACTGAGGC

HBV H- TATCAAGGAATTCTGCCCGTTTGTCCT

## Comment

The use of a nested approach facilitates sequencing of low level HBV DNA such as is found in the anti-HBe seropositive individual. The four sequencing reactions allow secure sequence data for assembly of the small HBs contig.

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