Diagnostic Testing and Interpretation of Lab tests for HBV

Brian Lam, MSHS, PA-C Inova Fairfax Hospital Center for Liver Diseases

Hepatitis B Nomenclature and/or Lab Tests (1)

- **HBV**: Hepatitis B virus.
- **HBsAg**: Hepatitis B surface antigen. Marker of infectivity when found in serum.
- **anti-HBs**: Antibody to HBsAg. Marker of immunity when found in serum.
- **HBcAg**: Hepatitis B core antigen. No commercial test available for this.
- anti-HBc: Antibody HBcAg. Marker of past or current infection.

Hepatitis B Nomenclature and/or Lab Tests (2)

•	gM anti-HBc : IgM is an antibody subclass of anti-HBc.
	Indicates recent infection with HBV
	(<4-6 mos.).

- **IgG anti-HBc**: IgG is a subclass of anti-HBc. Indicates "older" infection with HBV.
- **HBeAg**: Hepatitis B "e" antigen. Can only be present if HBsAg is positive. Marker of high degree of infectivity.
- Anti-HBe: Antibody to "e" antigen. May be present in infected or immune person.

Interpretation of Hepatitis B Panel

Hbs Ag	Hbc Ab	Hbs Ab	
-	-	-	Susceptible
-	+	+	Immune due to natural infection
-	-	+	Immune due to vaccination
+	Hbc Ab + HbcAb Igm +	-	Acutely infected
+	Hbc Ab + HbcAb Igm -	-	Chronically infected
-	+	-	4 possible interpretations

Four possible interpretations of isolated antiHBc positive

- **1.** May be recovering from acute HBV infection.
- **2.** May be distantly immune and test not sensitive enough to detect very low level of anti-HBs in serum.
- **3.** May be susceptible with a false positive anti-HBc.
- **4.** May be undetectable level of HBsAg present in the serum and the person is actually a carrier.

Tests for Measuring Clinical Response

Measuring Clinical Response

- Biochemical measurement of response
 - Liver enzymes
 - Normalization of ALT and AST
 - Synthetic tests
 - Normalization of albumin, bilirubin and INR
- <u>Serological measurement of response</u>
 - Loss of HbeAg +/- Gain of anti-HBe (HBeAg seroconversion)
 - Loss of HbsAg +/- Gain of anti-HBs (HBsAg seroconversion)
- Virologic measurement of response
 - HBV DNA suppression
- Histological measurement of response
 - Decrease in inflammation and fibrosis

Lok AS et al. Gastroenterology. 2001;120:1828-1853.

Why Focus on HBV Serum DNA?

- Serum HBV DNA is the best available measure of HBV replication
- HBV replication = disease
 - Reduction in HBV DNA precedes biochemical and histological response
 - Rebound in HBV DNA precedes loss of biochemical and histological response
 - Level of HBV replication correlates with incidence of HCC and cirrhosis

HBV DNA and Effect on Natural History

Risk for HCC and Cirrhosis

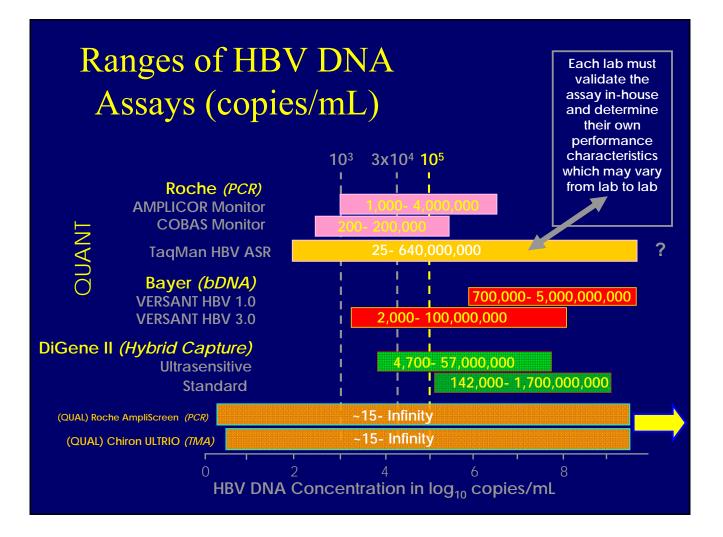
Assays for measuring HBV DNA

Nucleic Acid Tests for HBV DNA

- Signal amplification assays
 - Liquid hybridization assay (Abbott)
 - DNA-RNA
 - Branched DNA, bDNA (Bayer)
 - Hybrid capture (Digene)
- Target amplification assays
 - Polymerase chain reaction assay, PCR
 - Transcription-mediated amplification, TMA (Chiron)
 - Nucleic acid sequence-based amplification, NASBA
 - Ligase chain reaction assay, LCR

Lack of Standardization in HBV DNA Quantification

- Dynamic ranges of detection
 - Lower Limit of Quantification (LLQ)
 - Upper Limit of Quantification (ULQ)
- Presentation of results
 - Mean change in serum HBV DNA as percent reduction in HBV DNA
 - Log change is best format vs linear
 - Comparison vs. baseline
- Units of measurements
 - Number of units to express serum levels of HBV DNA
 - IU is now the standard



HBV DNA Evaluation

- HBV DNA should be evaluated using the assay with the widest dynamic range
- Patient who are negative by a quantitative assay should then be evaluated with the most (qualitative) sensitive assay available
- All patients should be treated until HBV DNA negative by the best assay available

HBV DNA and Effect on Natural History

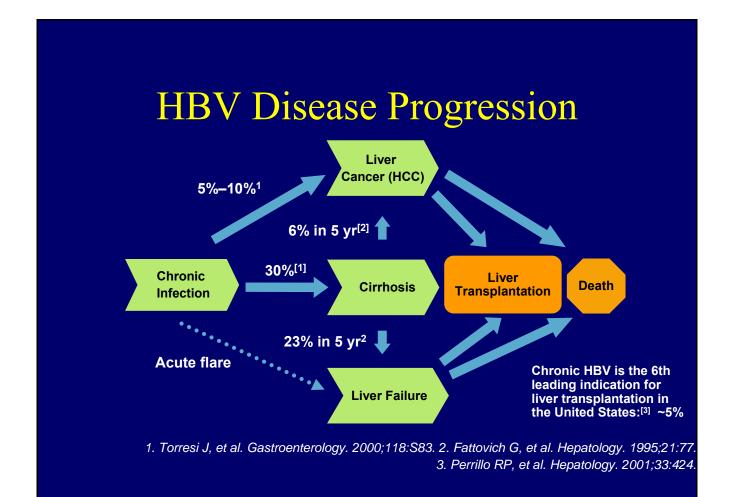
Risk for HCC and Cirrhosis

Factors Associated With Increased Risks of Progression to Cirrhosis

Host Factors	Virus Factors	Environmental Factors
Older age (longer duration)*	High levels of HBV replication*	Concurrent infection (HCV*, HDV, HIV)
Male*	Genotype $(C > B)^*$	Alcohol consumption*
Immune status	HBV variant (core promoter)	Diabetes mellitus [†] Obesity [†]

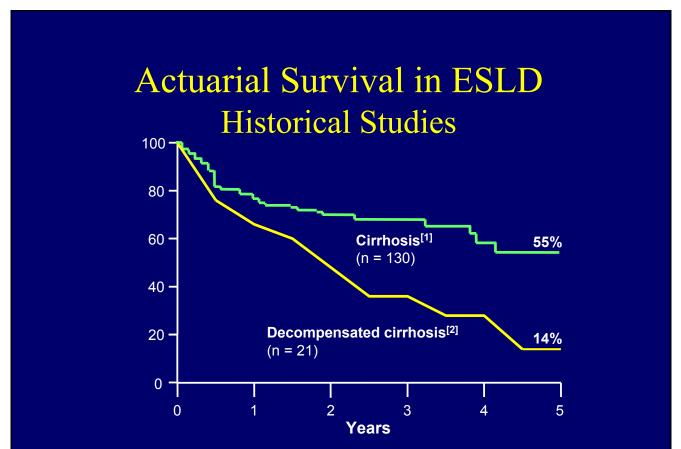
*Supported by strong evidence. [†]Further studies needed.

Yim, Lok. Hepatology 2006;43:s173-s181



Probability of Decompensation in Patients with Compensated Hepatitis B Cirrhosis

N = 349100 Mean f/u period 80 of 73 months 60 % At 5 years, 40 probability of decompensation 20 was 23% 0 3 5 6 7 8 9 10 2 4 0 Year Fattovich G, et al., Hepatology 1995; 21:77

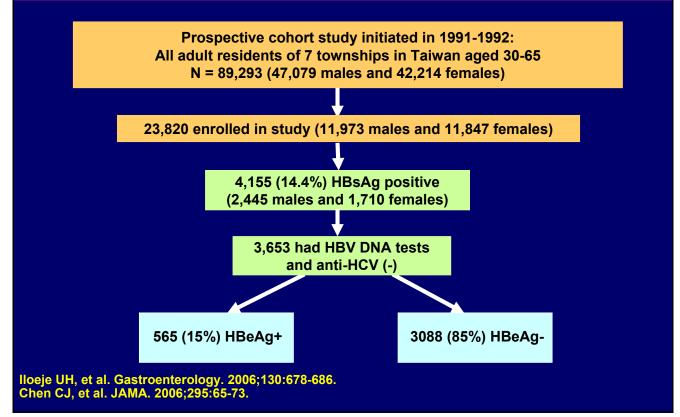


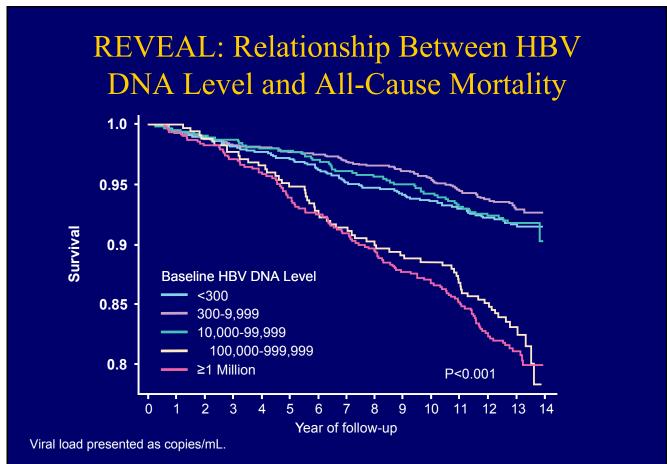
1. Weissberg JI, et al. Ann Intern Med. 1984;101:613. 2. De Jongh FE, et al. Gastroenterology. 1992;103:1630.

REVEAL

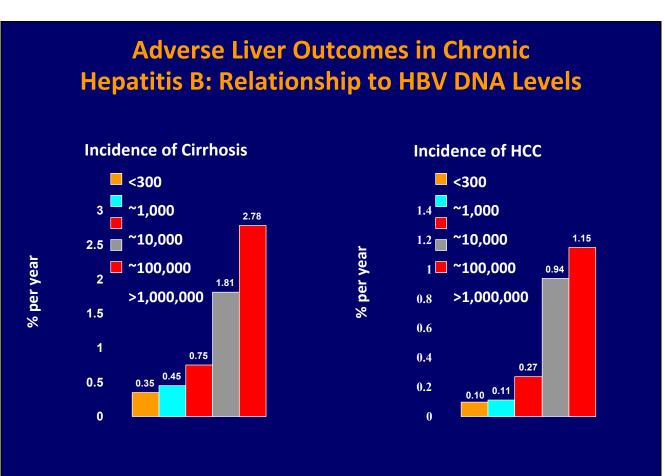
Taiwan Cohort Study



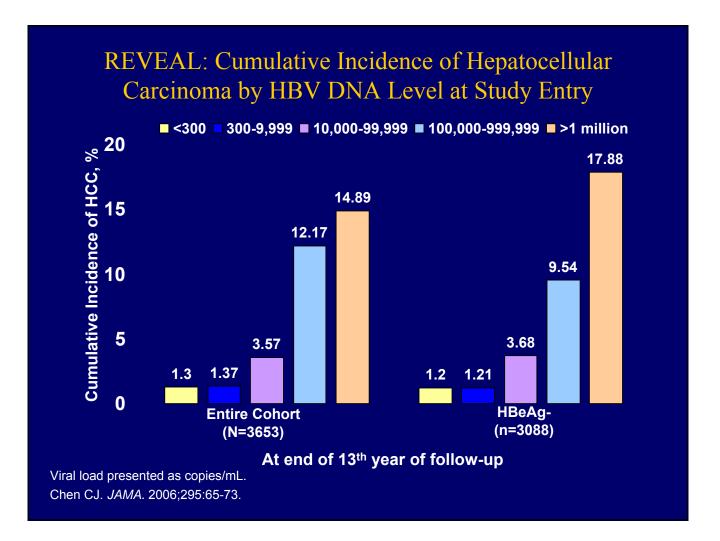




Iloeje UH. Gastroenterology. 2006;130:678-686.



1. lloeje UH et al. *Gastroenterology* 2006;130:678-686. 2. Chen C-J et al. *JAMA* 2006;295:65-73.



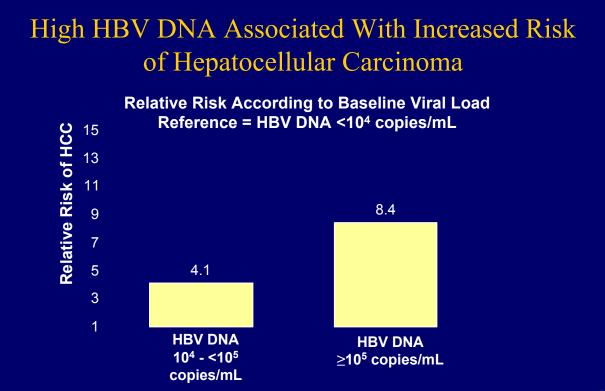
REVEAL: Relationship Between Persistent Viremia and HCC Incidence

• Persistent HBV DNA associated with greater risk of HCC

- Examined in those with levels $\geq 10^4$ copies/mL at baseline (n = 1376)

1.2 x 10 ⁴ 1.0 x 10 ⁴ 1.0 x 10 ³ 3.0 x 10 ³ 4.0 x 10 ³ 4.0 x 10 ³ 2.0 x 10 ³ 0 4.0 x 10 ³	1473	5882	8730	10,108
Baseline HBV DNA, copies/mL	< 10 ⁴	≥ 10 ⁵	≥ 10 ⁵	≥ 10 ⁵
Follow-up HBV DNA, copies/mL		< 10 ⁴	10^4 to < 10^5	≥ 10 ⁵
Adjusted RR (95% CI)	1.0 (reference)	3.8 (1.7-8.4)	7.3 (3.5-15.3)	10.1 (6.3-16.2)
<i>P</i> Value len CJ, et al. <i>JAIMA</i> 2006;295:65-	 •/3.	< .001	< .001	< .001

Cł



Nested case-control series. 37 cases matched with 61 controls. Median viral load was 9.3×10^5 copies/mL in the HCC cases and undetectable (<3x10⁴ copies/ml) in the controls (*P*<0.001).

Evans AA. 55th AASLD, October 29 - November 2, 2004, Boston, MA.

Impact of Viral Replication on Disease Progression: Taiwan Cohort Study Results

- The incidence of hepatocellular carcinoma (HCC) and liver cirrhosis is correlated with level of viral replication
- Persistent elevation of viral load over time has the greatest impact on HCC risk
- Viral load predicts risk of future HCC independent of HBeAg status and serum ALT level
- This risk increases with increasing viral load

EASL April 2005

HBV Genotype?

HBV Genotype

Currently eight genotypes (A-H) defined

Clinical Outcomes

- Genotype B: better than genotype C
- Genotype A: better than genotype D
- Genotypes E to H: the clinical significance remains to be examined

Sustainted response to standard interferon

- Genotype B better than genotype C
- Genotype A better than genotype D.

Response to PegIFN

• A and B, better rates of HBeAg seroconversion than C and D

<u>Responses to nucleos(t)ide analogues</u>

• No significant difference between HBV genotypes have been identified

HBV Genotype Conclusion

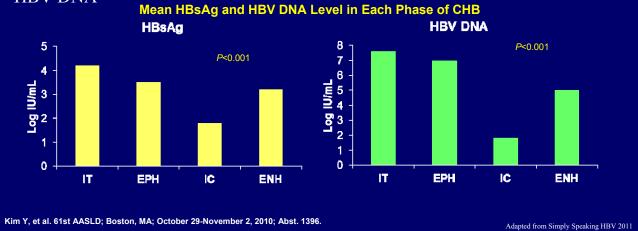
Currently

- HBV genotyping is not routine
- but this is likely to change if additional studies confirm that genotyping has prognostic value or strongly predicts response to therapy

HBsAg levels?

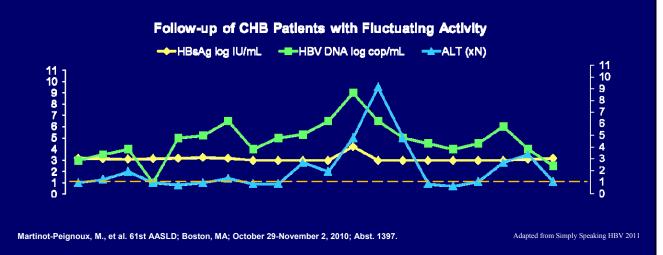
HBsAg Levels in the Natural Course of HBV

- HBsAg levels have been recently studied as markers of response to treatment
- Study evaluating HBsAg level variance with stage of disease
- Cross sectional retrospective study, N = 601
- HBsAg correlates with HBV DNA except in HBeAg(-) patients
- In untreated patients, higher HBsAg levels may predict recurrence as well as HBV DNA



Quantitative HBsAg: New Marker for Diagnosis of HBsAg Inactive Carriage

- Quantitative HBsAg is new marker for diagnosis of HBV inactive carriage
- HBsAg ≤2,000 IU/mL and HBV DNA ≤2,000 copies/mL identifies true inactive carriers with 100% specificity
- Yearly measurement of HBsAg to identify patients that will demonstrate HBsAg seroconversion
- Conclusion: Quantitative HBsAg should be considered for future HBV guidelines



HBsAg Levels in the Natural Course of HBV

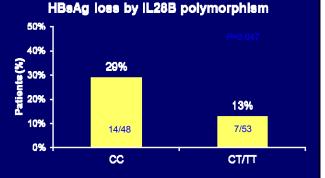
Key Message:

- HBsAg levels are known to correlate with treatment success with interferon.
- This study demonstrates that they parallel HBV DNA and can be used to predict the course of HBV infection, even when not on treatment.
- measurement of HBsAg to identify patients that will demonstrate HBsAg seroconversion
- Conclusion: There may be clinical utility in following HBsAg levels

IL28B Testing?

IL28B Polymorphism May Predict HBsAg Clearance in Genotype D, HBeAg(-) Patients Treated with Interferon Alfa

- Retrospective study of role of IL28B polymorphism in interferon treatment in HBeAg(-) CHB pts
 - Treatment: standard IFN for 23 months, 11 year follow-up
 - Endpoint HBsAg loss
- Low baseline HBV DNA, high ALT levels and genotype CC of IL28B independently predicted HBsAg clearance
- IL 28B polymorphism may represent an additional pretreatment predictor of interferon response in HBeAg(-), genotype D patients with CHB



Predictors of HBsAg Loss by Multivariate Analysis

Factors	OR	(95% CI)	P value
HBV DNA levels <6 log cp/mL	9.9	(2.5 - 38.7)	0.001
ALT levels >136 IU/mL	5.3	(1.5 -17.9)	0.007
IL28B genotype CC	3.9	(1.2 -12.4)	0.023

Lampertico P, et al. 61st AASLD; Boston, MA; October 29-November 2, 2010; Abst. 237.

Adapted from Simply Speaking HBV 2011

Treatment Options







Natural Clearance of HBeAg

- Occurs in 4% to 12% of carriers per year
- 40% to 50% of HBeAg+ carriers will clear HBeAg in 5 years
- 70% to 80% will clear HBeAg in 10 years
- Occurs more frequently in older carriers and those with elevated aminotransferase levels
- 20% of carriers who clear HBeAg will have one or more reversions to HBeAg+

Alaska HBV Carrier Study

McMahon Ann Int Med 2001;135:759-68

- Population based cohort study: all ages, genders
- 1536 HBV carriers tested semiannually
- F/u 19,430 person years (average 12.3)
- 641 initially HBeAg-positive
- Probability of HBeAg clearance 73% by 10 years

Clearance of HBsAg

- Occurs in ~ 0.5% of carriers per year
- Occurs in 10% to 15% of carriers who clear HBeAg with interferon therapy
- No evidence HBsAg clearance occurs more frequently in carriers who clear HBeAg during Lamivudine therapy



HBV Natural History

Progression to cirrhosis and decompensation



REVEAL

HBeAg Status and HCC

