## Diagnostics for Control of Hepatitis A What do we Need and Why?

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- Role of Diagnostics in Vaccine Preventable Diseases
- Hepatitis A Diagnostics
- Specialized Assays for Hepatitis A
  - -Serologic
  - -Molecular



# Some "Definitions"

- Diagnostics
  - -widely used
  - -commercially available
- Specialized Assays
  - Generally for research only
  - Not widely available
  - May be commercially available



## Diagnostic Tests and Assays: Vaccine Preventable Diseases

#### Pre - Vaccine

Diagnosis of acute disease (epidemiology / disease burden)

#### Vaccine Assessment

- Detection of infection
- Assessment of vaccine response
- Vaccine Introduction
  - Effectiveness
  - Long-term effects



# **Events in Hepatitis A Virus Infection**





Adapted from: J Med Virol 1988; 26:315-326; J Infect Dis 2000; 182:12-17

## **Diagnosis of Acute Disease**

- Differential diagnosis of jaundice and acute febrile illness
- Clinical management
- Surveillance
  - Outbreak detection
  - Disease burden estimates
  - Post-introduction vaccine effectiveness
- Epidemiologic studies
- Clinical trials



# **IgM Anti-HAV**

- An excellent <u>diagnostic</u> test among persons with symptoms suggestive of hepatitis
  - High sensitivity & specificity
  - High predictive values positive and negative
  - "Detuned" to improve specificity only positive 4-6 months after symptom onset
- Transiently positive following vaccination (8-20%) – usually not a diagnostic problem

# IgM Anti-HAV

- The *downside* = not widely used in countries where hepatitis A is endemic
  - differential diagnosis of acute hepatitis
  - (IgM anti-HAV & IgM anti-HBc)
  - non-icteric syndromes that could be hepatitis A (e.g., febrile illness in children)
  - Relatively high cost
  - No rapid test formats

## **Assessment of Hepatitis A Vaccination**

#### Short-term Vaccine Response (total anti-HAV)

- Clinical trials
- Epidemiologic studies
- Problems:
  - Diagnostic test = lower levels of detection
  - Diagnostic test must be modified, not generally applicable to vaccinated persons
  - Measures antibody to structural proteins (vaccine and wild-type infection)



## **Assessment of Hepatitis A Vaccination**

#### Long-term

- Antibody persistence (total anti-HAV)
- Breakthrough infections
  - Clinically evident (IgM anti-HAV)
  - Inapparent problematic
    - Virus detection have to be lucky
    - Antibody to HAV non-structural proteins



## **Assessment of Hepatitis A Vaccination**

- Antibody to non-structural (replication) antigens of HAV
  - Response to proteins produced during viral replication
  - Not present following vaccination with inactivated viral vaccines
  - Could identify subclinical infections in vaccinated population



## Studies of Antibodies to HAV Non-Structural Proteins



#### **CDC Group**

Robertson BR, et al J Med Virol 176: 593 (1993) **NIH Group** 

Stewart DR, et al JID 176: 593 (1997) Kabrane-Laziz Y, et al. Vaccine 19: 2878 (2001)

## **Antibodies to Non-Structural Proteins**

- Proof of concept: antibodies can be detected
- Limitations = sensitivity
  - High viral replication = high rate of detection (>95%)
  - –Low viral replication (e.g., attenuated vaccine) = low rate of detection (~25%)
  - Poor detection of persons with low levels of viral replication (small sample sizes)
  - —Unknown identification of persons with breakthrough infections following vaccination



## Summary Serologic Antibody Assays / Tests

Excellent diagnostic test – IgM anti-HAV -More widespread use Possible need for special assays -Total anti-HAV more sensitive -Antibody to non-structural proteins (anti-C3) more sensitive



# **Molecular Diagnostics**



ses

Virus Detection

 Humans during infection
 Environmental samples

 Molecular epidemiology

 Transmission patterns
 Virus evolution



# **Events in Hepatitis A Virus Infection**





Adapted from: J Med Virol 1988; 26:315-326; J Infect Dis 2000; 182:12-17

## **Detection of HAV RNA in Serum**

#### Time from symptom onset to blood draw

| Days    | Positive (%) |  |
|---------|--------------|--|
| <0      | 100          |  |
| 0 -13   | 93.4         |  |
| 14 -27  | 93.5         |  |
| 28 - 41 | 63.3         |  |

#### Not affected by source of infection, gender, race, or age



Source: J Infect Dis 2000; 182:12-17 and CDC unpublished data

## Virus Detection in Environmental Samples Challenges

- Material often NOT same material implicated in outbreak
- Foods (e.g., berries, onions, shellfish)
  - Special extraction methods to release virus from food surfaces / matrices and large biomass
  - Concentration of extracts from large volumes
- Water and sewerage
  - Large volumes require concentration (e.g., membranes)

Multiplex for other enterically transmitted agents (e.g., noro and caliciviruses)

## Virus Detection in Environmental Samples

#### Nucleic acid amplification

- Inhibitors from food components, or elution and concentration methods
- Detection of infectious virus
  - Immuno-capture RT-PCR
- –Amplification methods
  - Dependent on throughput needs and lab capacity (e.g., real time, quantitative, RT-PCR)



## Regions Commonly Used to Amplify Hepatitis A Virus



#### From: Clinical Microbiology Reviews (2006) 19: 63

# **Genetic Relatedness of HAV**



5-15

- Relatively low degree of nucleotide variation across genome regions
- 7 genotypes

– 4 human

- 3 simian
- Enough variation to determine relatedness of isolates using relatively short sequence fragments

# **Uses of Molecular Epidemiology**

Sources of Virus Transmission

 Food / water / other environmental
 Risk factors – MSM, IDU
 Blood / Blood Products

 Transmission Patterns within Populations
 Monitoring Vaccine Effectiveness



# **Sources of Virus Transmission**

# Food / water / other environmental Simultaneous outbreaks in multiple locations Multiple food sources – e.g., berries, green onions, shellfish

## Risk factors

- -Outbreaks in IDUs
- Disease transmission patterns among MSM
- Transmission Patterns after Vaccination



## Multistate Outbreak of Hepatitis A Associated with Frozen Strawberries, 1997

### Lessons Learned

- Could identify small number of cases using markers of genetic relatedness
- Required high-throughput molecular diagnostics
- Required large data base of genetic sequences for general population
- Required previously agreed upon sequenced regions for comparison



Hutin et al. NEJM 1999; 340:595-602

## **Relatedness of HAV from Cases who Ate Frozen Strawberries from Same Processor**

| State      | # Cases | # Sera<br>available | # with outbreak sequence |
|------------|---------|---------------------|--------------------------|
| Michigan   | 198     | 118                 | 118                      |
| Tennessee  | 2       | 1                   | 0                        |
| Wisconsin  | 5       | 5                   | 5                        |
| Louisiana* | 4       | 2                   | 2                        |
| Maine      | 29      | 10                  | 8                        |
| Arizona    | 10      | 7                   | 7                        |
| USA        | _       | 98                  | 4                        |



\*Commercial product

#### Multi-state Outbreak of Hepatitis A Associated with Frozen Strawberries, United States, 1997



Sentinel Counties Tacoma, Portland, Birmingham

Michigan (118) Louisiana (2) Maine (8) Arizona (7) Wisconsin (5)

Hutin et al. NEJM 1999; 340:595-602

# Summary

- Have powerful tools for molecular diagnostics
- Genetic markers (molecular epidemiology) has increased our knowledge of HAV transmission
- Must continue sharing information about strains
- We have tools to show elimination of HAV transmission in immunized populations

Vaccines don't Prevent Disease Vaccination Prevents Disease



# **Dedication**

## **Omana Nainan**

## **Betty Robertson**

