



# Challenges in diagnosis of Hepatitis E virus infections

Suzan D. Pas

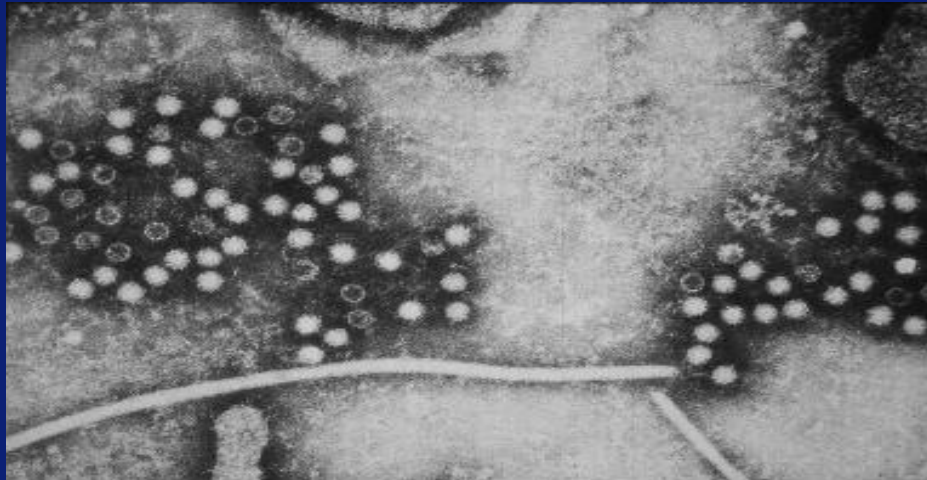
Afdeling Virologie

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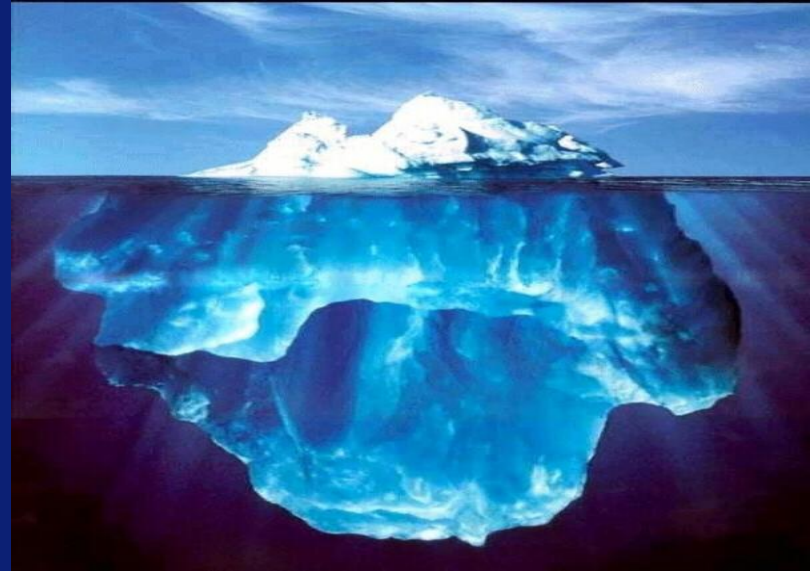
# Hepatitis E Virus

- Hepeviridae family
- Non-enveloped virus
- Positive sense, single stranded RNA of  $\pm 7200$  bp
- 27-34 nm in diameter

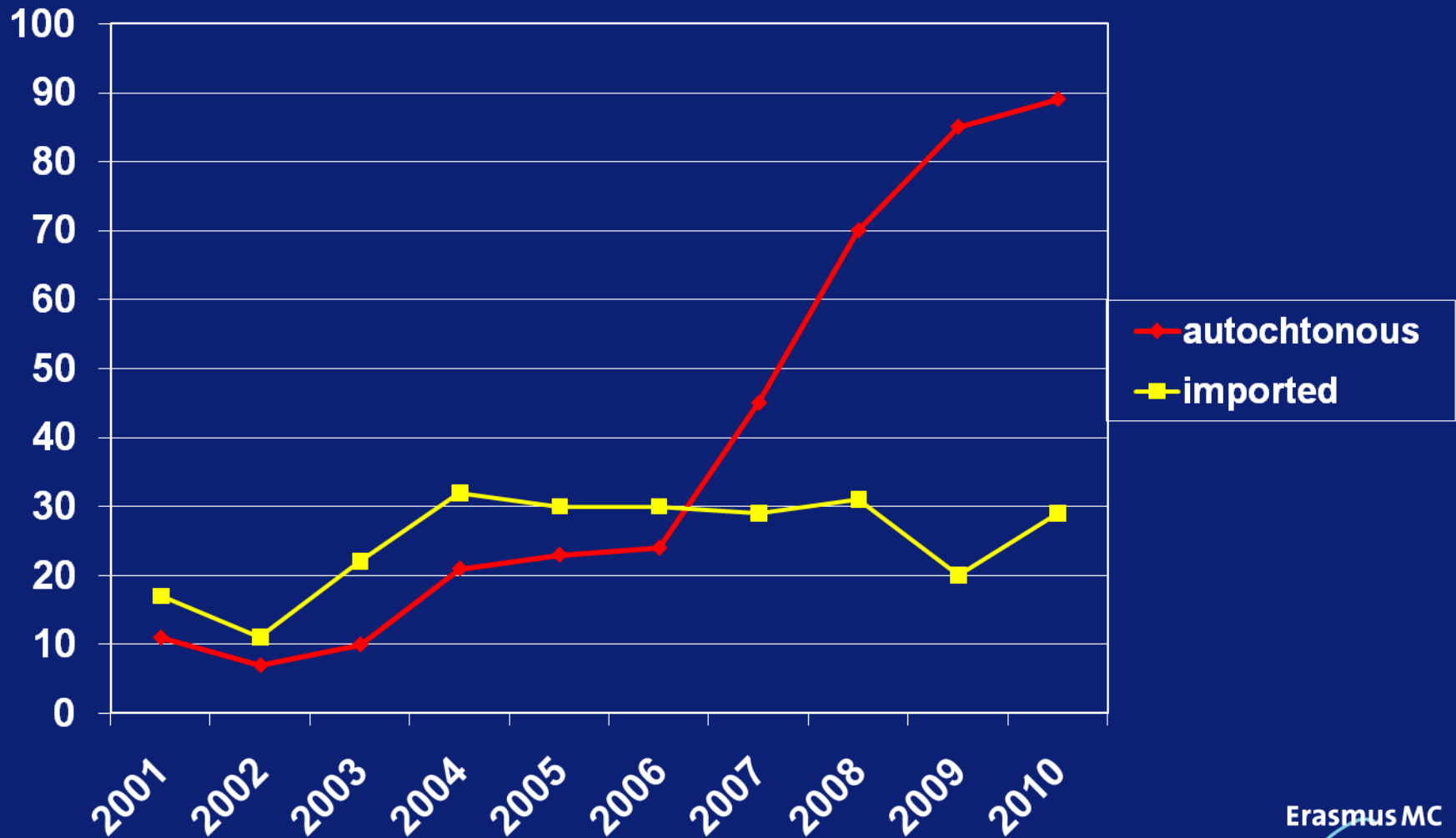


# Clinical presentation of hepatitis E virus

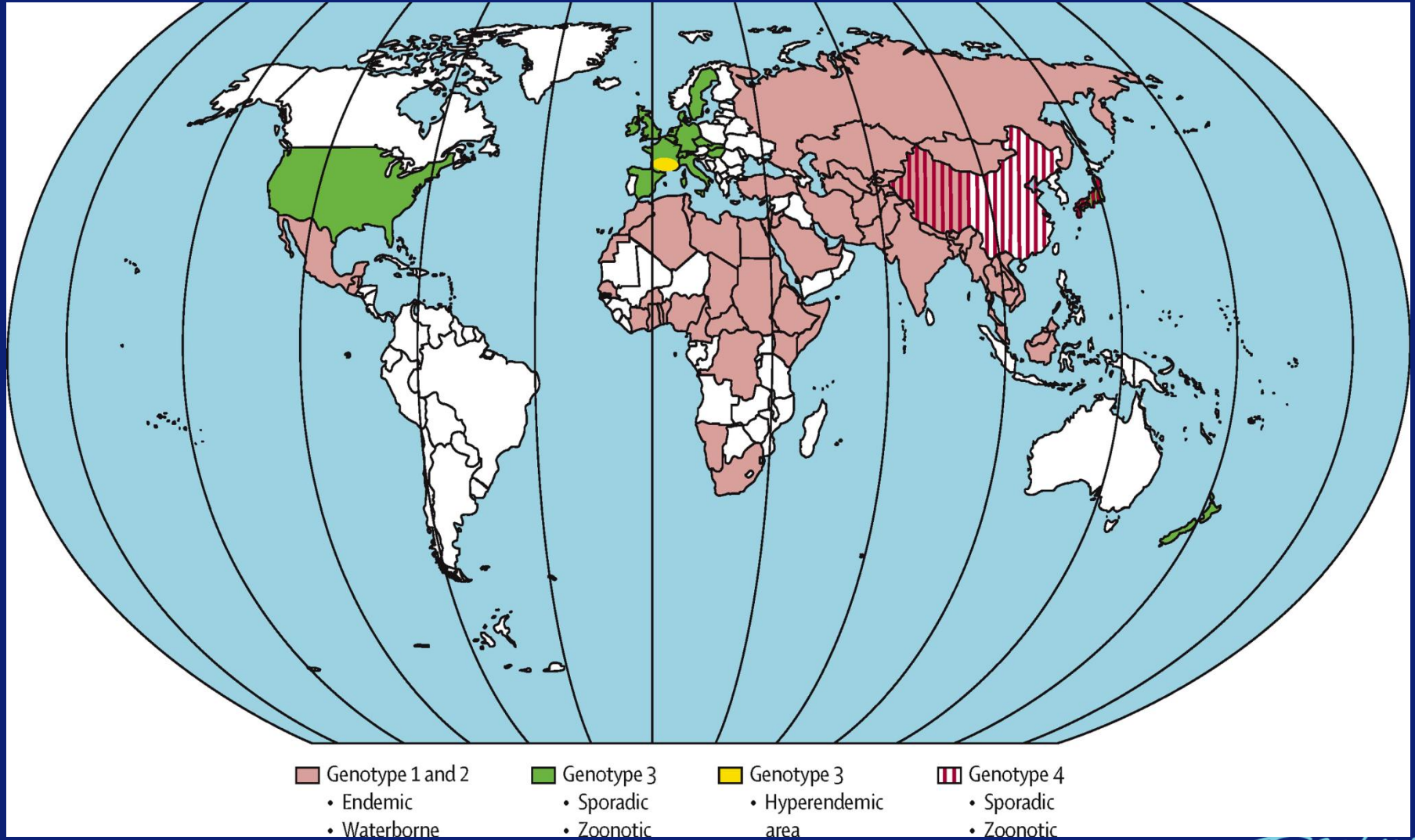
- Fever
- Fatigue
- Loss of appetite
- Nausea
- Vomiting
- Abdominal pain
- Jaundice
- Dark urine
- Clay-colored stool
- The ratio of symptomatic to asymptomatic infection is reported to range from 1:2 to 1:13.
- Mortality: overall 1-4%, pregnant women 15-25%



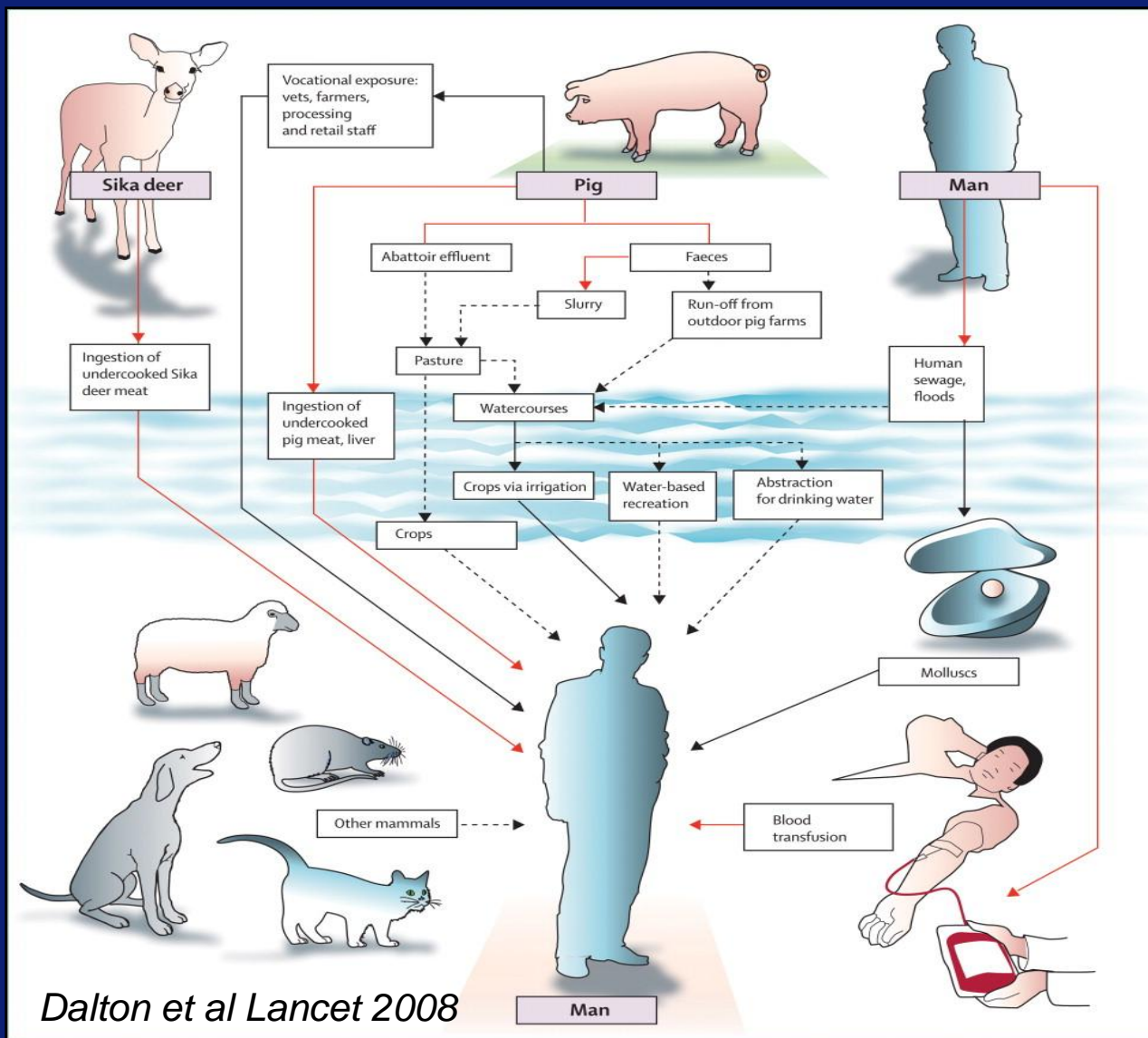
# Number of reported HEV-cases in Germany



# Global distribution of HEV genotypes



# HEV transmission in developed countries - zoonose

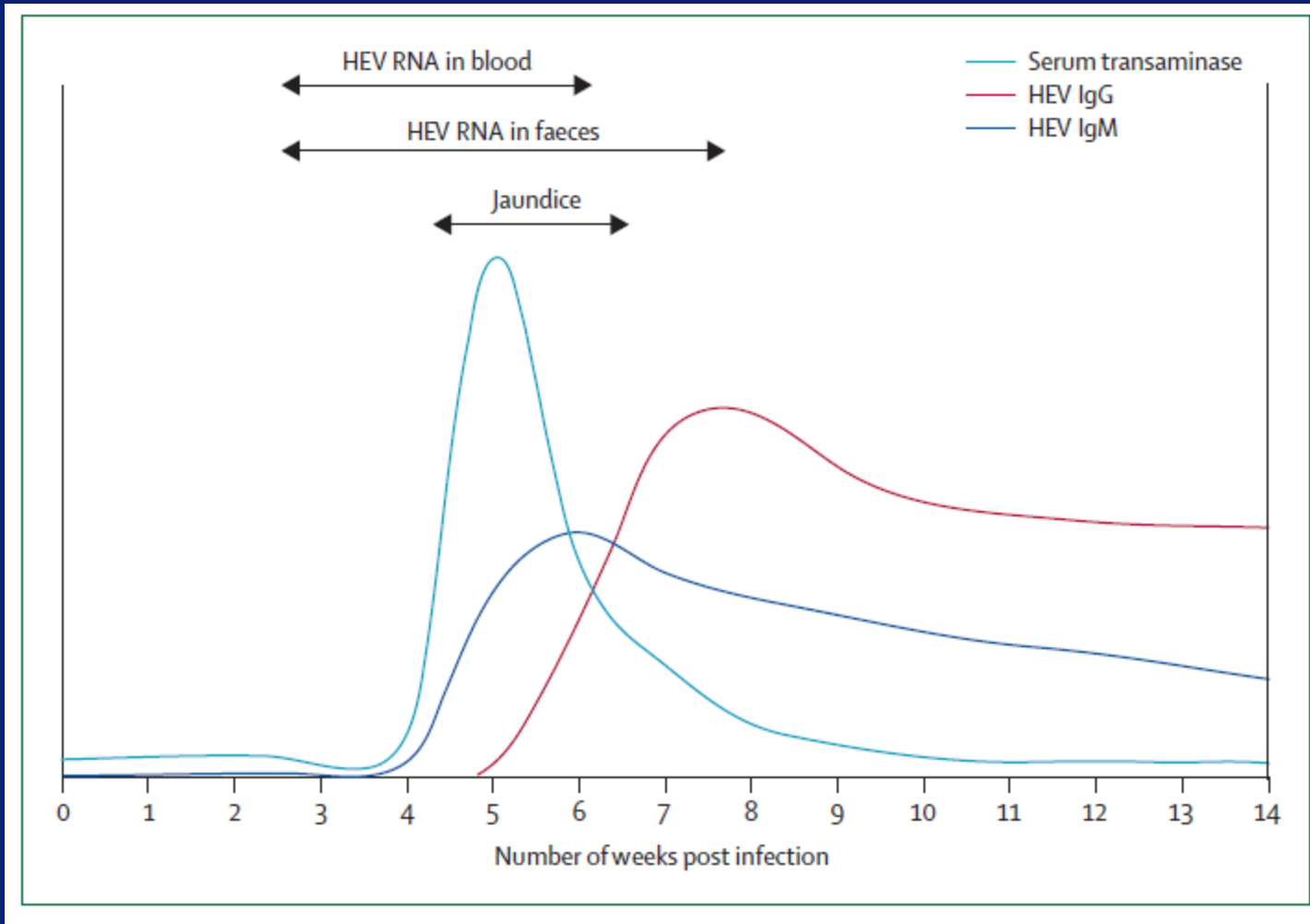


*~55% of fecal tanks in Dutch pig farms HEV RNA positive EID 2007*

*~ 6% of Dutch porcine livers HEV RNA positive. J Food Prot. 2007*

Dalton et al Lancet 2008

# Course of HEV infection in the immunocompetent



Dalton et al, Lancet 2008

# Chronic HEV in solid organ recipients

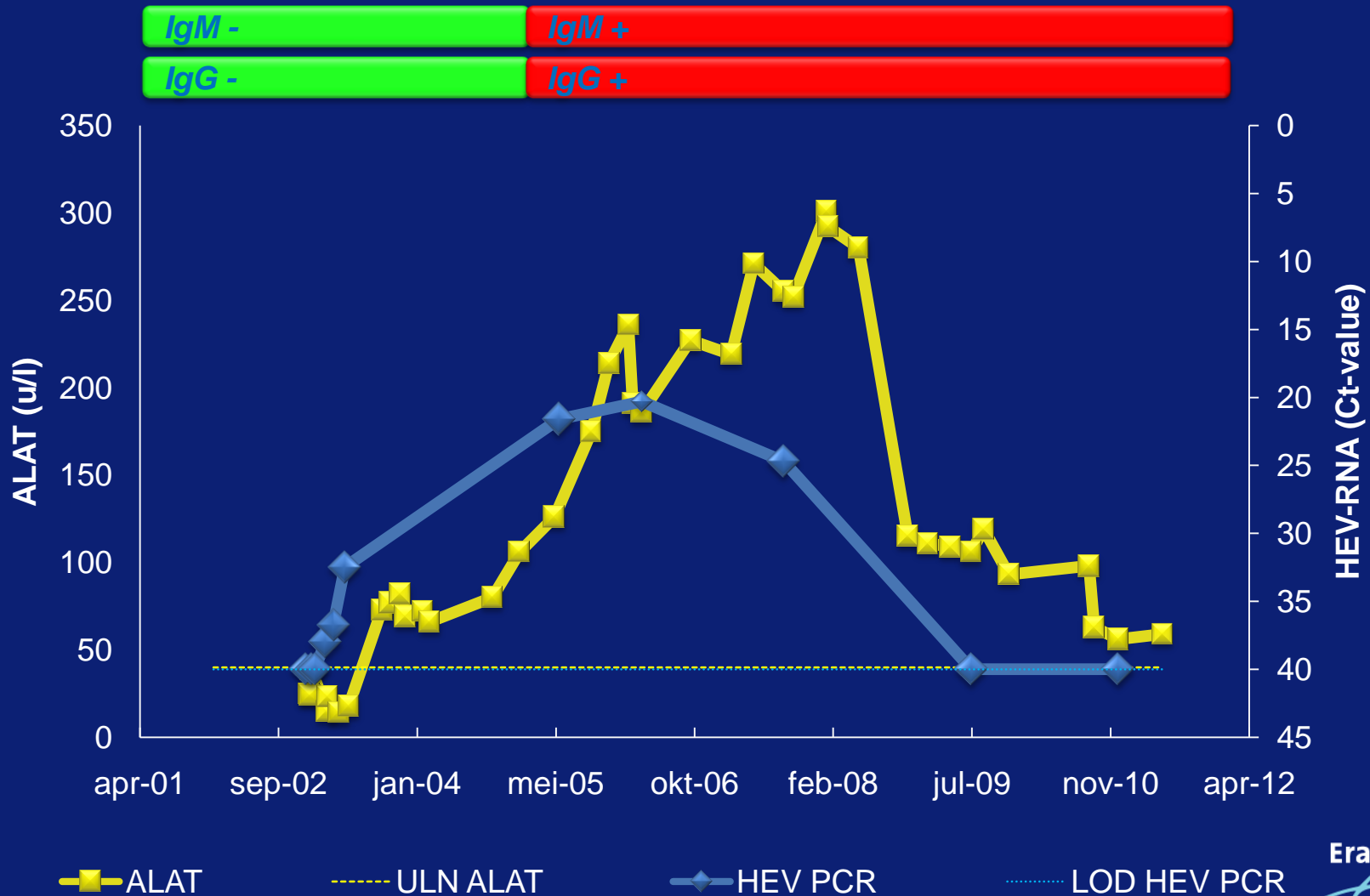
- Chronic HEV infection reported in the transplant setting
  - Persistent viraemia
  - Persistently raised transaminase activity
  - Histological features associated with chronic hepatitis
  - Evidence of rapid development of cirrhosis
- Association with a more profound immunosuppression

Kamar, N Engl J Med 2008;358(8):811-7.

Haagsma, Liver Transplantation 2009;15(10):1225-8

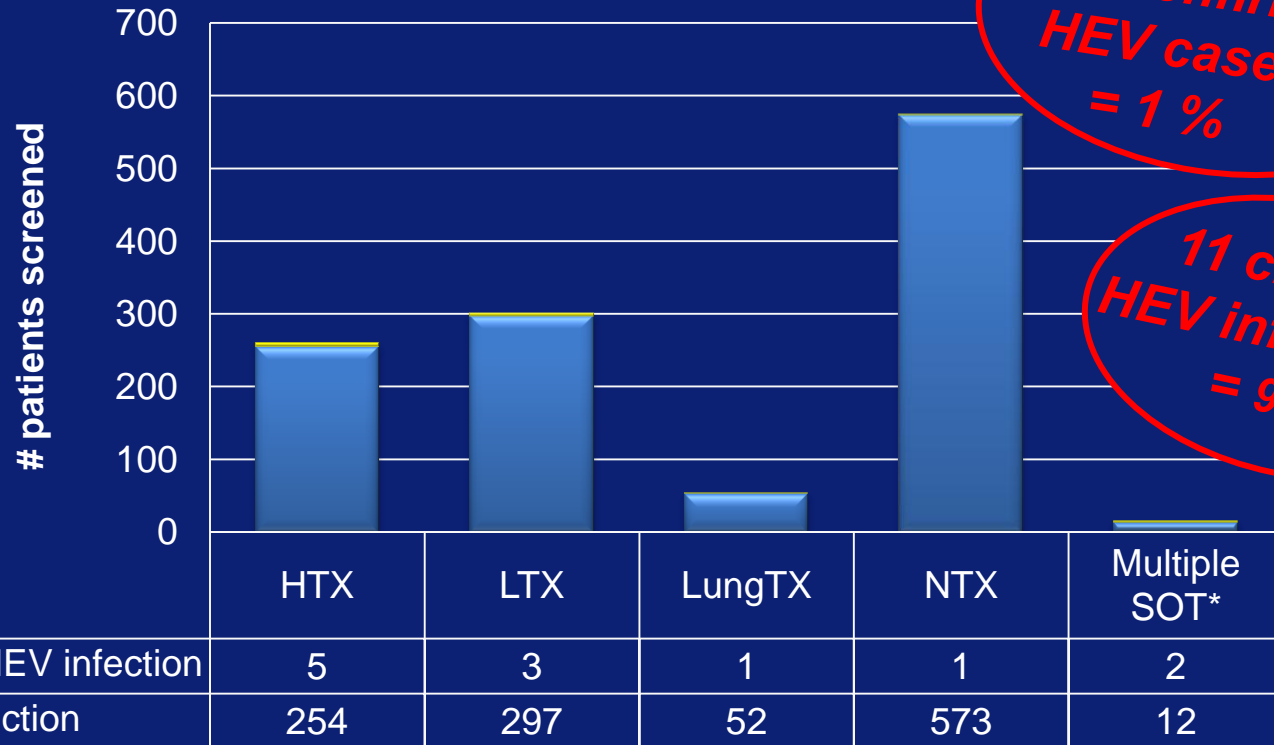


# Chronic HEV infection misdiagnosed as Graft v Host



# HEV in living adult SOT transplant recipients

## HEV infections among SOT patients



12 confirmed HEV cases = 1 %

11 chronic HEV infections! = 91,6 %

Mixed Tx group:

confirmed HEV case: 1 NTX-LTX , 1 NTX-HTX

no HEV infection: 8 NTX-LTX, 3 NTX-HTX, 1 NTX-LuTX

Erasmus MC



# Current status of HEV diagnostics

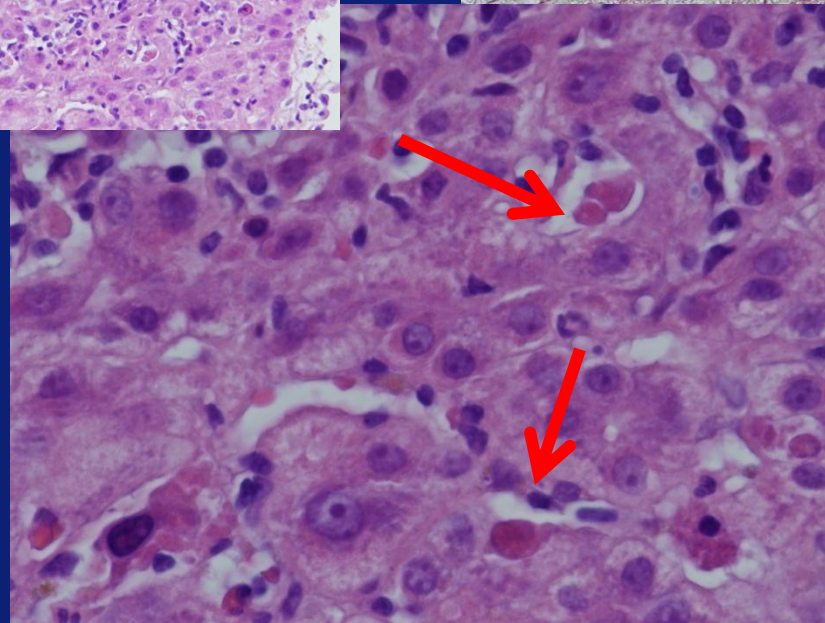
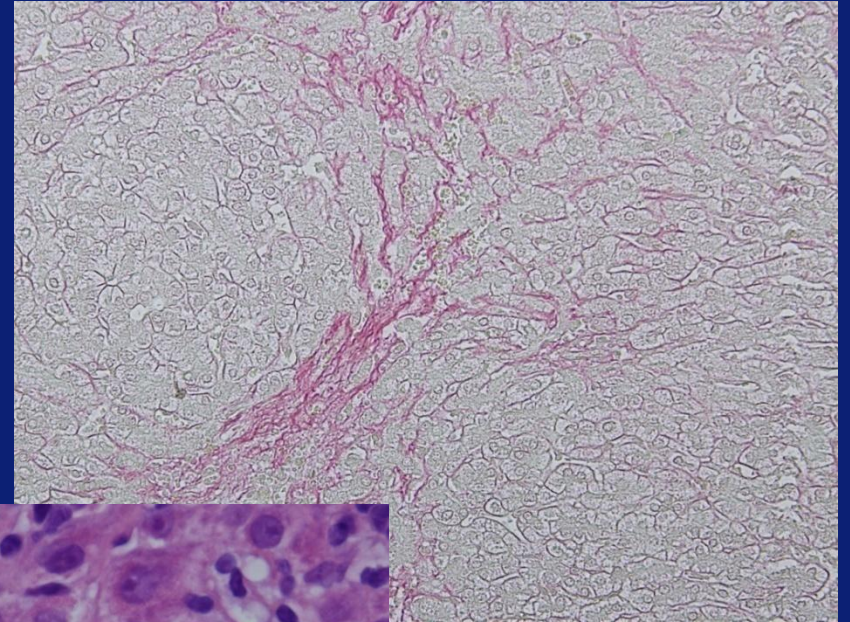
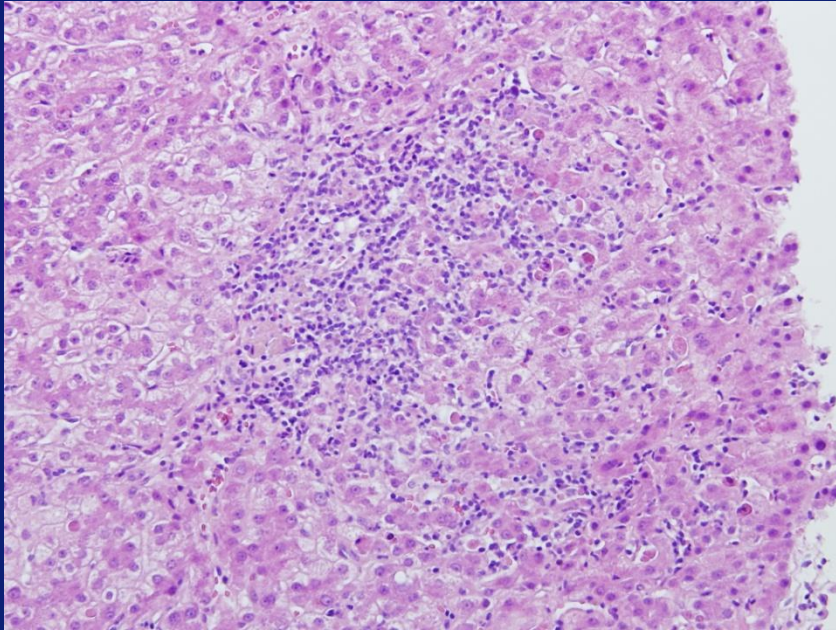
\* **Pathology**            not specific  
                                 Invasive

# Current status of HEV diagnostics - Histopathology



Liver biopsy overview

# Current status of HEV diagnostics - Histopathology



# Current status of HEV diagnostics

- \* Pathology    not specific  
                  Invasive
  
- \* **Virus culture** - inefficient

# Virus culture -HEV

Journal of Medical Virology

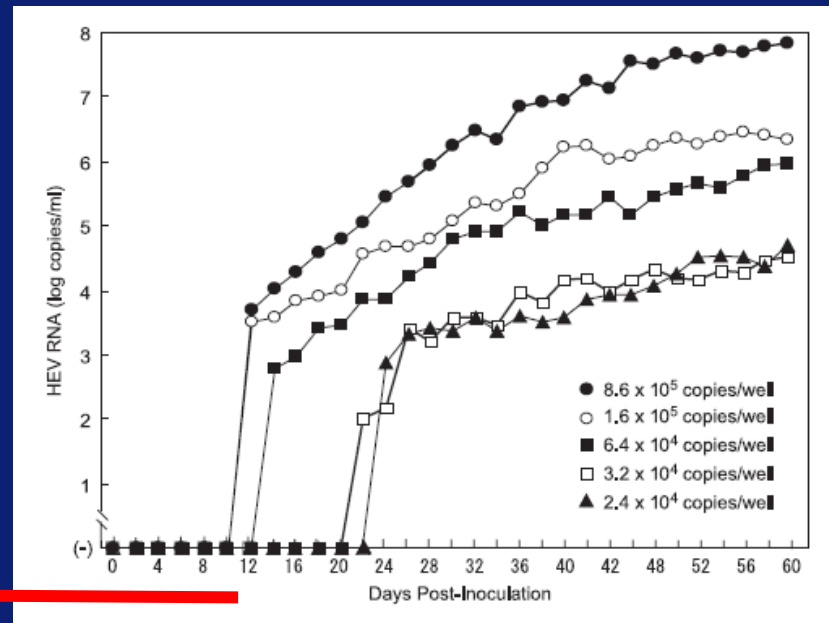
*Rev. Med. Virol.* 2011; 21: 18–31.  
Published online in Wiley Online Library  
(wileyonlinelibrary.com)  
DOI: 10.1002/rmv.678

## Efficient cell culture systems for hepatitis E virus strains in feces and circulating blood

Hiroaki Okamoto\*

*Division of Virology, Department of Infection and Immunity, Jichi Medical University School of Medicine, Shimotsuke-Shi, Tochigi, Japan*

PLC/PRF/5 (hepatocellular carcinoma) and A549 (lung cancer) cells



*long incubation  
period, low  
sensitivity*



# Current status of HEV diagnostics

- \* Pathology not specific

  - Invasive

- \* Virus culture - inefficient

- \* **HEV serology**

  - validation of commercial assay

  - conformational testing using blot



# HEV serology - literature

*A lot of in house assays described*

CLINICAL AND VACCINE IMMUNOLOGY, May 2007, p. 562–568  
1556-6811/07/\$08.00+0 doi:10.1128/CVI.00231-06  
Copyright © 2007, American Society for Microbiology. All Rights Reserved.

Vol. 14, No. 5

## Use of Serological Assays for Diagnosis of Hepatitis E Virus Genotype 1 and 3 Infections in a Setting of Low Endemicity<sup>▽</sup>

M. Herremans,\* J. Bakker, E. Duizer, H. Vennema, and M. P. G. Koopmans

*IgM and IgG* **Genelabs**, *Mikrogen Recomblot*

Journal of Medical Virology 82:799–805 (2010)

## **A Comparison of Two Commercially Available Anti-HEV IgG Kits and a Re-Evaluation of Anti-HEV IgG Seroprevalence Data in Developed Countries**

Richard Bendall,<sup>1\*</sup> Vic Ellis,<sup>1</sup> Samreen Ijaz,<sup>2</sup> Rachel Ali,<sup>3</sup> and Harry Dalton<sup>3</sup>

**IgG** ; *Genelabs, Wantai, 4.5x higher seroprevalance with Wantai*

Rasmus MC

*Ezafung*

# HEV serology - literature

## Serologic Assays Specific to Immunoglobulin M Antibodies against Hepatitis E Virus: Pangenotypic Evaluation of Performances

Jan Drobeniuc,<sup>1</sup> Jihong Meng,<sup>1,2</sup> Gábor Reuter,<sup>3</sup> Tracy Greene-Montfort,<sup>1</sup> Natasha Khudyakova,<sup>1</sup> Zoya Dimitrova,<sup>1</sup> Saleem Kamili,<sup>1</sup> and Chong-Gee Teo<sup>1</sup>

CID 2010:51 (1 August) •

*2 in-house assay, 4 commercial available assays*

	Sens.(%)	Spec.(%)
Int. Immuno-Diagnostics	82	91,2
MP biomedical (former Genelabs)	72	93
RPC Diagnostic Systems	98	95
Mikrogen	92	95,6

*Sens. panel 50 samples, 4 genotypes*

*Spec. panel 229 samples*

# HEV Serology validation - included ELISAs

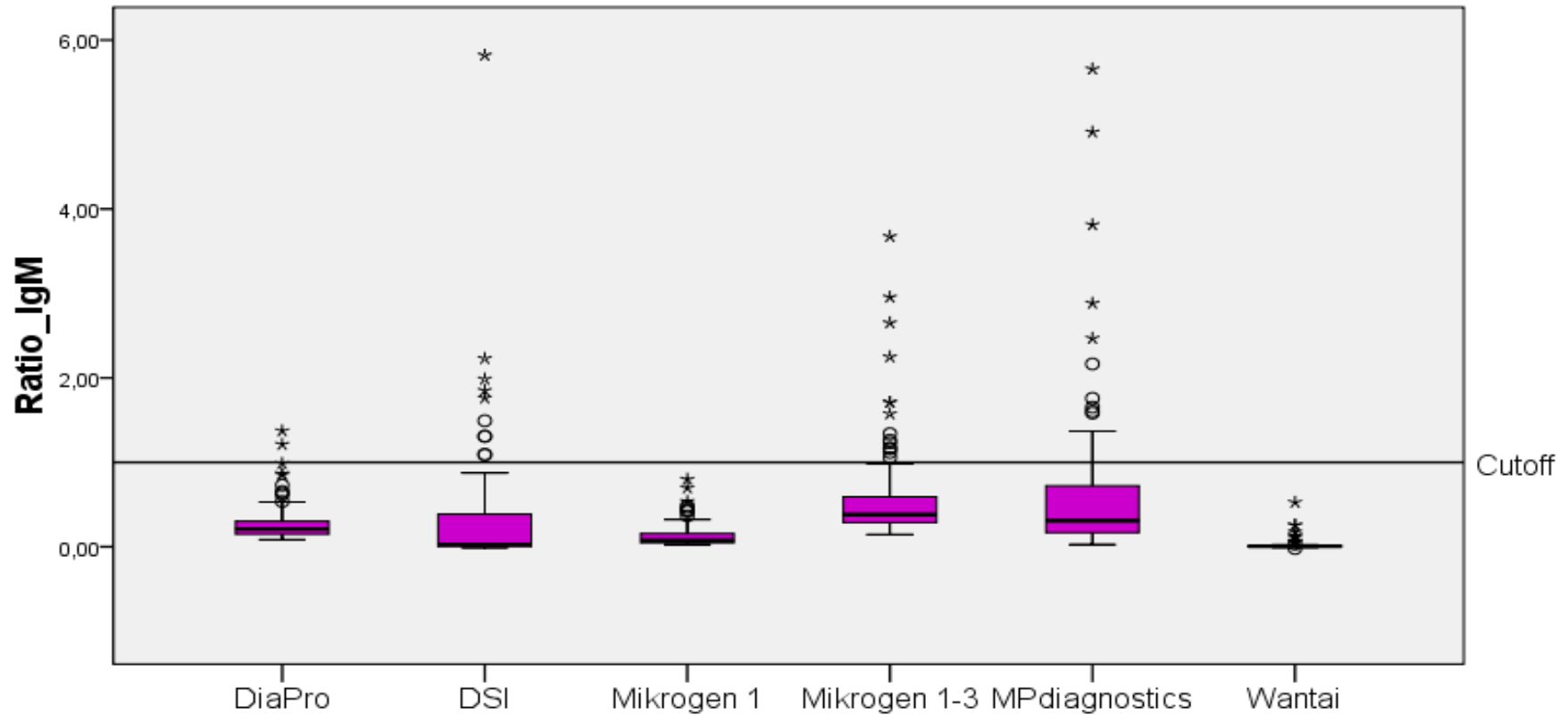
	Name	Company	Country	Genotype inclusivity	antigens
IgM/ IgG	HEV Elisa v3.0	MP diagnostics	Singapore	gt 1 en 2	mix of peptides from ORF2 and complete ORF3
	<i>recomWell</i> HEV IgM/IgG	Mikrogen Diagnostik	Germany	gt 1	synthetic, ORF2 en ORF3 (e.coli)
	<i>recomWell</i> HEV IgM/IgG <b>new</b>	Mikrogen Diagnostik	Germany	gt 1 and 3	synthetic, ORF2 en ORF3 (e.coli)
	HEV IgM/IgG	DRG	Germany	gt 1 en 2	synthetic, ORF2 en ORF3
	HEV IgM/IgG	Dia.Pro	Italy	gt 1 en 2	synthetic, ORF2 en ORF3
	HEV IgM/IgG	RPC Diagnostic systems / <b>DSI</b>	Italy	gt 1 en 2	artificial ag. Composed of 12 antigenic regions derived from ORF2 and ORF 3
	HEV IgM/IgG	DiaCheck			
	HEV IgM/IgG	Wantai HEV IgG PE2	Singapore		PE2 peptide from structural region of ORF2

# Sensitivity panel : HEV PCR confirmed patients

Time of Drawal	# samples	# patients	# samples immune status			genotypes included		
			ID	non-ID	unknown	gt 1	gt 3	unknown
Prior to infection	12	12	12	0	0	0	9	3
< 6 wks	34	31	16	14	4	7	18	9
6 wks <t< 6 months	22	19	15	4	3	0	16	6
>6 months	20	16	16	3	1	1	17	2
<b>total</b>	<b>88</b>		<i>Division was made on basis of clinical symptoms in combination with retrospective HEV RT-PCR analysis</i>					

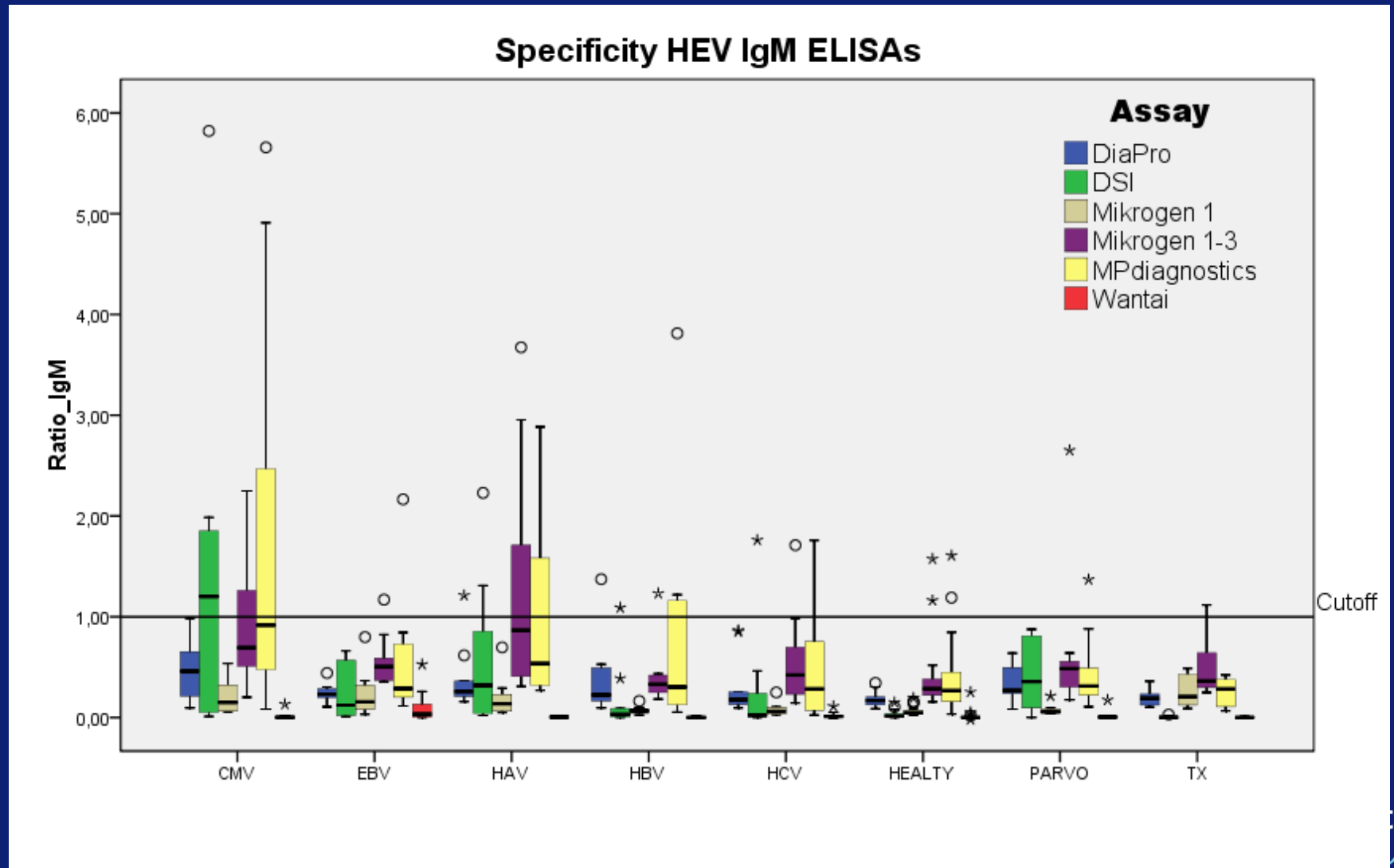
# Specificity panel HEV- IgM

## Specificity IgM ELISAs



*N= 89 samples*

# Specificity panel HEV- IgM



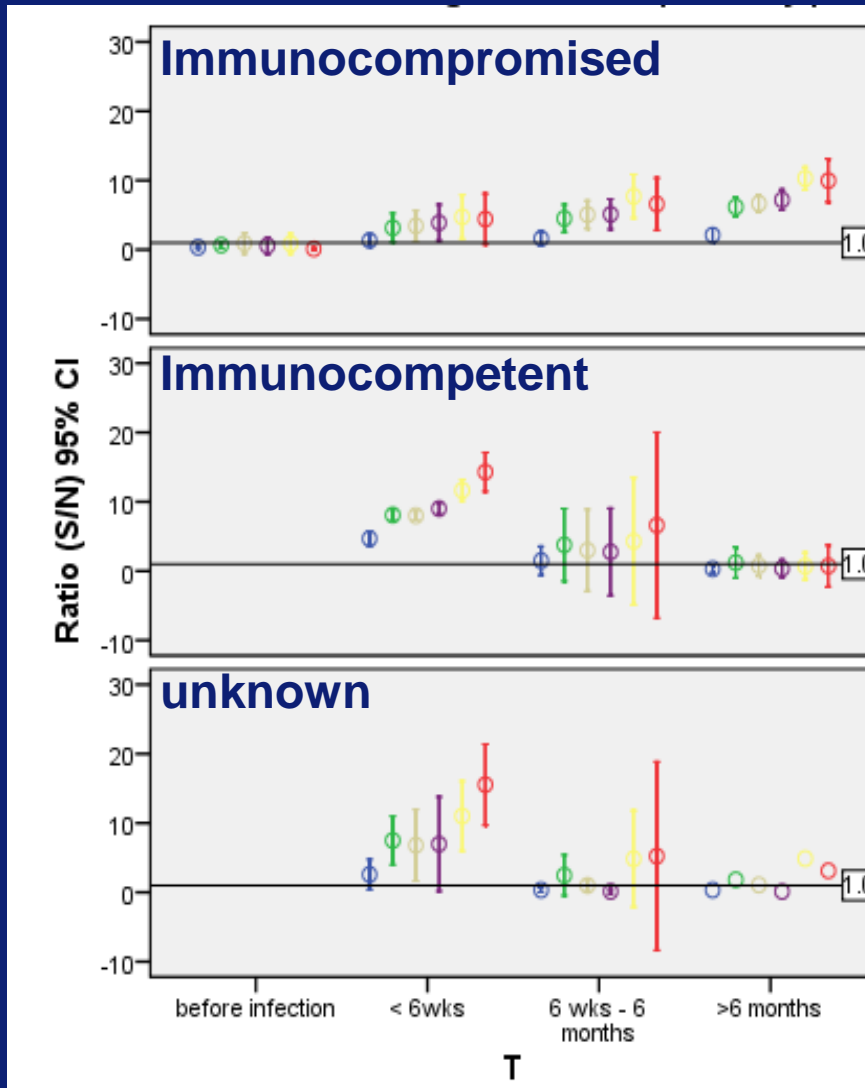
# HEV serology – validation of commercial assays

	Analytical sensitivity (titers at cutoff)					Clinical performance	
	IgM		IgG			Sens. IgM	Spec. IgM
	geno 1	geno 3	WHO	geno 1	geno 3		
Mikrogen old	4000	250	1600	6400	800	52%	>99%
Mikrogen new	32000	16000	3200	>12800	3200	79%	90%
MP diagnostics	>64000	4000	800	3200	100	74%	84%
DSI	8000	4000	800	3200	800	71%	90%
DiaPro	32000	32000	800	6400	100	81%	98%
Wantai	>64000	>64000	1600	>12800	800	75%	>99%
DRG	32000	32000	800	6400	100	excluded	
Diacheck	4000	250	400	3200	100		

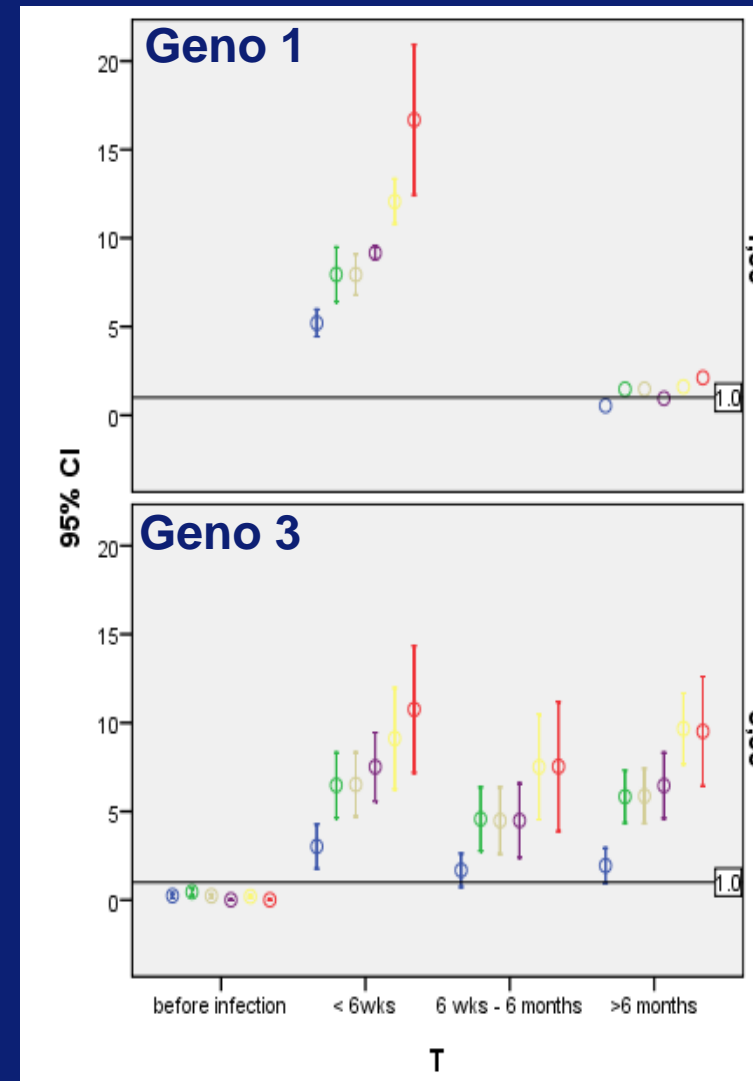
# IgM ratio's sensitivity panel

- Ratio\_Mikrogen\_1\_IgM
- Ratio\_Mikrogen\_13\_IgM
- Ratio\_MP\_IgM
- Ratio\_DSI\_IgM
- Ratio\_DiaPro\_IgM
- ratio\_wantai\_IgM

*per immune status group*



*per genotype*

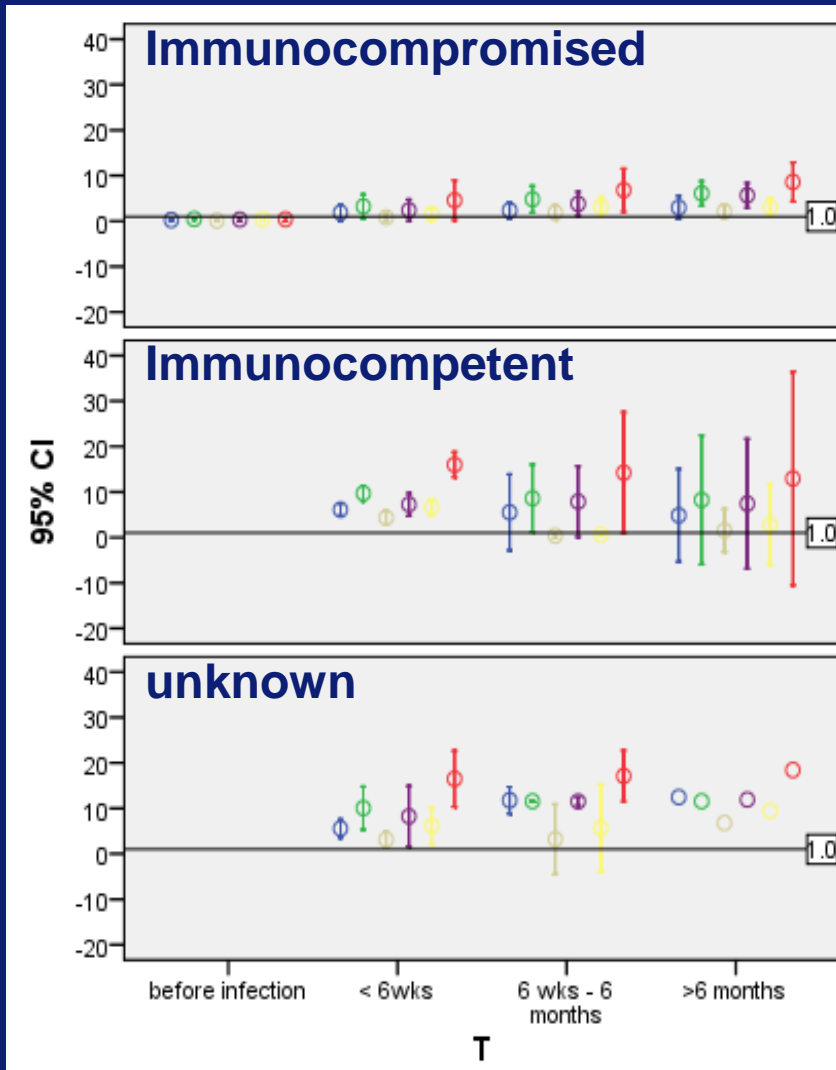




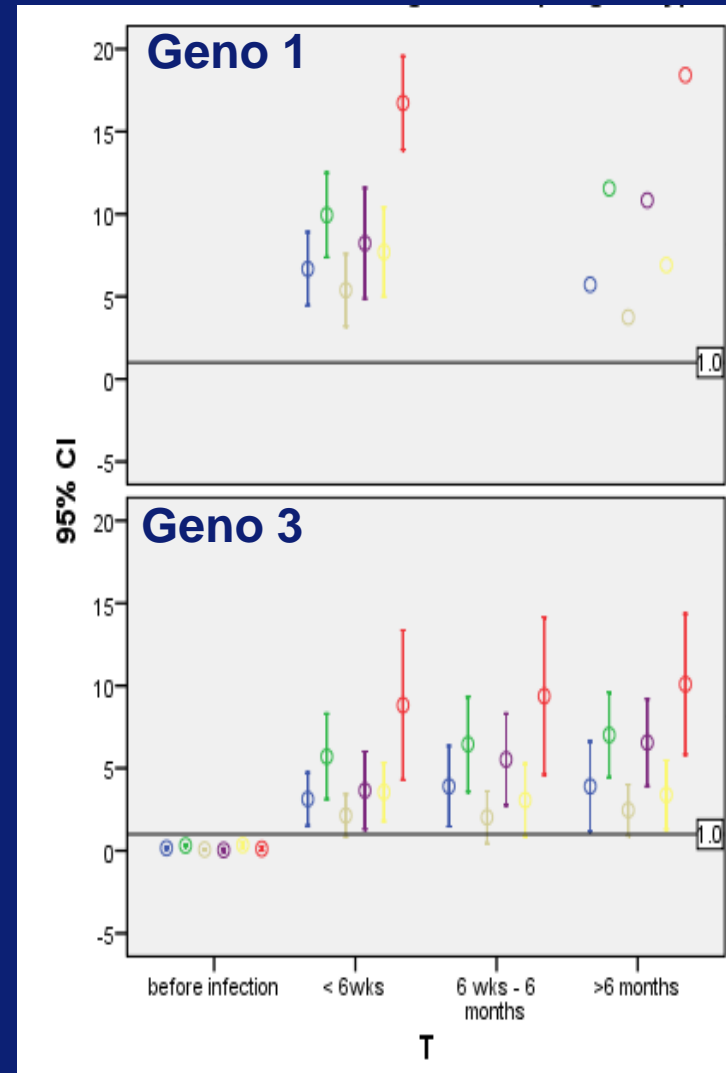
# IgG ratio's sensitivity panel

- Ratio\_Mikrogen\_1\_IgG
- Ratio\_Mikrogen\_13\_IgG
- Ratio\_MP\_IgG
- Ratio\_DSI\_IgG
- Ratio\_DiaPro\_IgG
- ratio\_wantai\_IgG

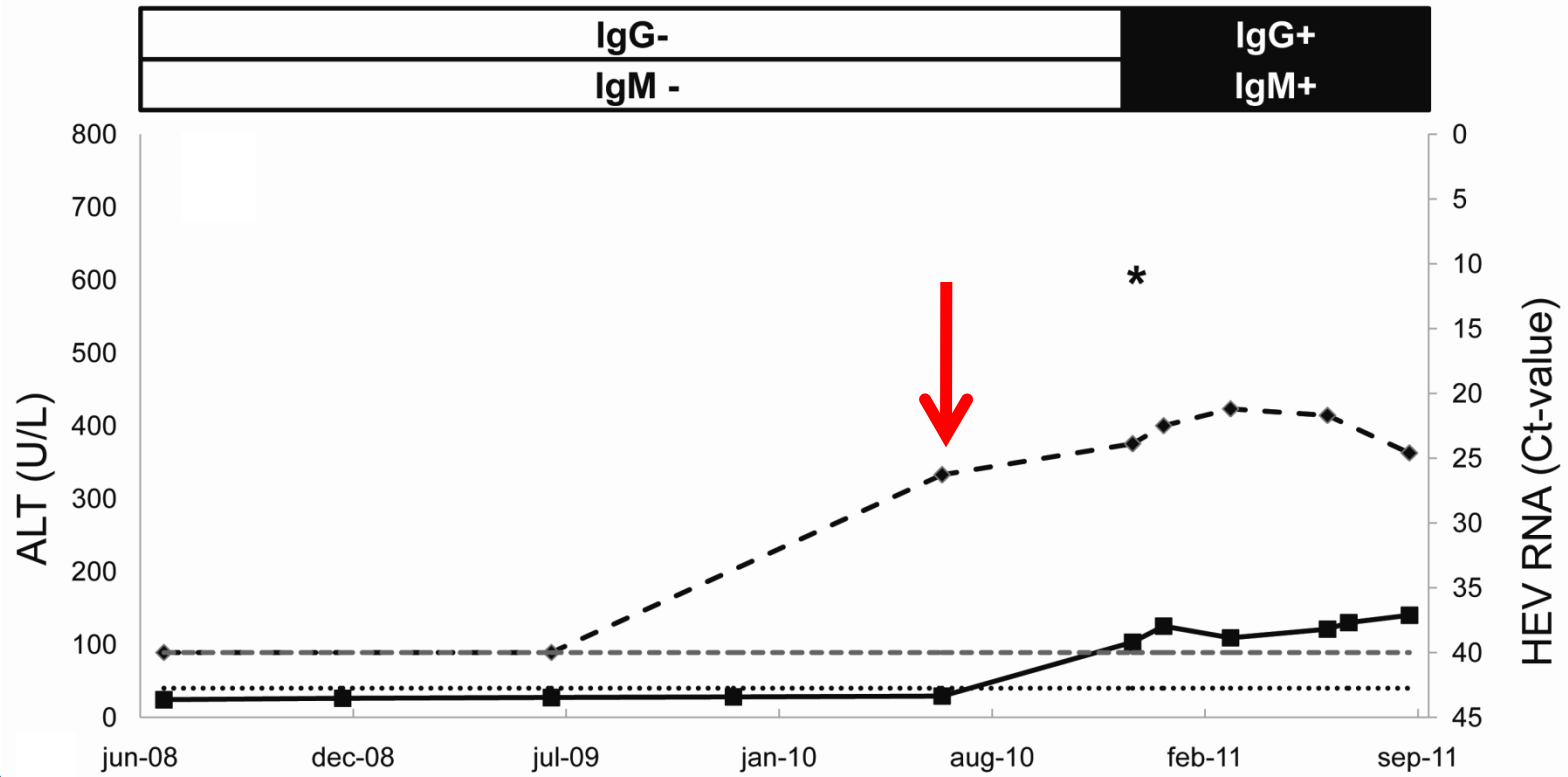
*per immune status group*



*per genotype*



# Course of chronic HEV infection



—■— ALT

-♦- HEV PCR

..... ULN ALT

---- LOD HEV PCR

## Parameters of HEV cases of SOT group

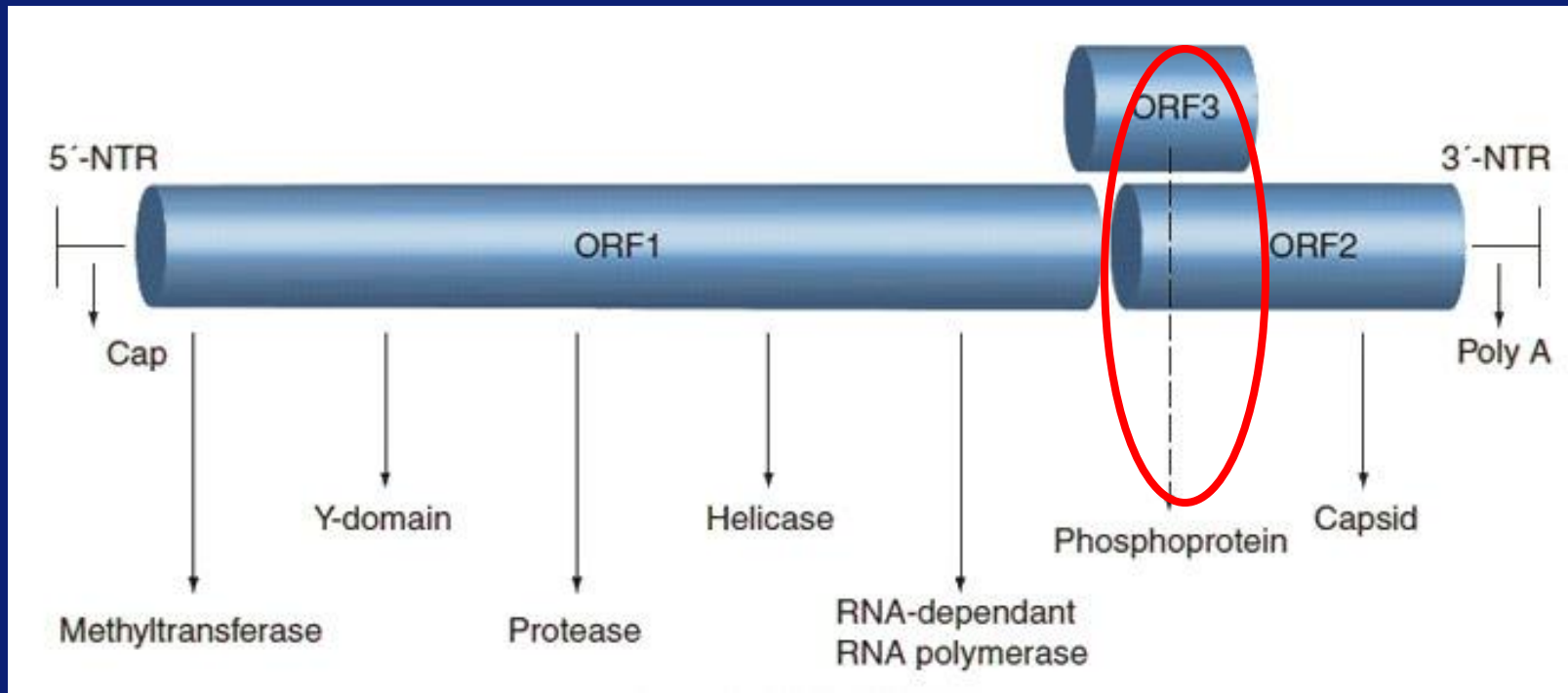
	median	range	ULN (F/M)
Peak ALT (U/L)	301	81 - 909	30/40
Peak AST (U/L)	172	66 - 1016	30/36
Peak $\gamma$ -GT (U/L)	299	72 - 1740	34/49
Peak Billirubine ( $\mu$ mol/l)	16	5 - 100	16/16
Peak HEV-RNA (Ct values)	20.0	16.7 - 26.6	NA
Period of HEV-RNA positivity (months)	16	6 - 55	NA
Time between the SOT and first HEV-RNA positive (months)	2.0	-0.3 - 20.1	NA
Time of HEV-RNA positivity prior to HEV IgM (days)	32	0 - 826	NA
Time of HEV-RNA positivity prior to HEV IgG (days)	124	0 - 826	NA

\* HEV IgM and IgG (Wantai Biochemicals)

# Current status of HEV diagnostics

- \* Pathology    not specific  
                  Invasive
- \* Virus culture    inefficient
- \* HEV serology
  - validation of commercial assay
  - conformational testing using blot
- \* **Molecular diagnostics**
  - **real time vs conventional RT-PCR**
  - **standardisation**
  - **genotyping**

# HEV real time RT-PCR



# HEV real time RT-PCR

1st Author	YoP	Validated	Target	Principle	Length	Limit of Detection
Mansuy et al	2004	Gt1 and 3		Two-step Taqman on Light cycler		1E3copies/ml
Orru et al	2004			SyBRgreen		10GEC
Jothukumar et al	2004	Gt1-4	ORF2/ORF3	taqman	70 bp	4GEC, others claim no quant.
Ahn et al	2006	gt3	ORF2/ORF3	taqman	103bp	16 copies/ml
Enouf et al	2006	Gt1-4	ORF2/ORF3	Taqman on Light cycler	86bp	10 copies/rx
Li et al	2006	GT 1-2, macaques	ORF2	SyBRgreen	207bp	4.5x10 <sup>3</sup> copies/rx
Gyarmati et al	2007	Gt1-4 2 human samples	ORF2	Taqman on Light cycler	113 bp	1-20 geq/rx
Zhao et al	2007	Gt1-4, theoretical	ORF2/ORF3	taqman	103bp	5,6E <sup>3</sup> copies/ml pseudovirus
Ward et al	2009	swine isolates only	comparison of four of the above ; Jothukumar et al most sensitive			
Vasichova et al	2012	swine isolates only	dual target Jothukuman2006+ Gyarmati2007	taqman in LC480	70/113bp	50 copies/ml



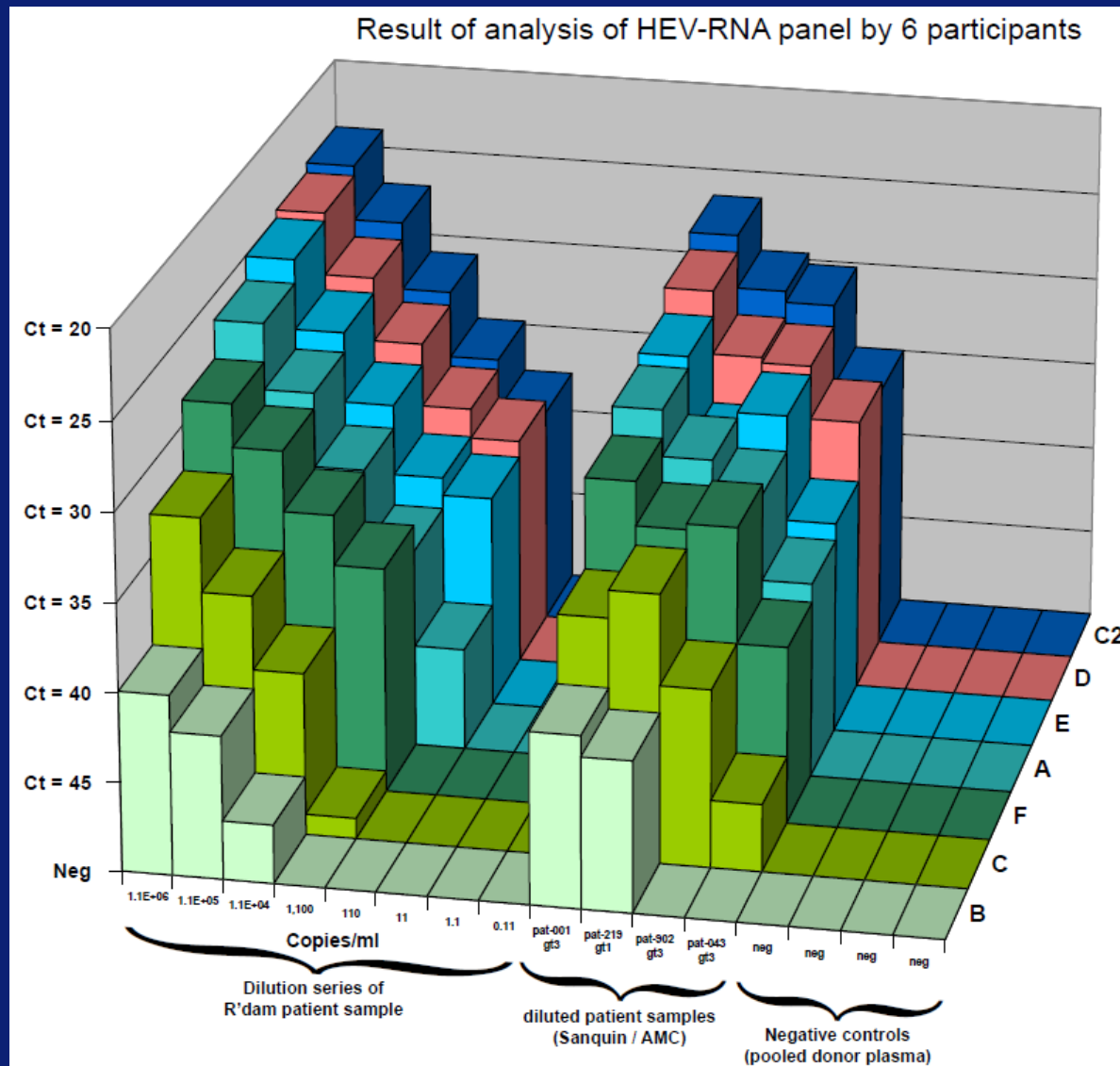
# HEV NAT quality assessment 2011 – Sanguin/ R'dam



- *6 participating labs of Dutch HEV workgroup*
- *Samples were randomized and included:*
  - *10 log dilution series of genotype 3 sample*
  - *calibrated against candidate WHO HEV standard\**
  - *four (diluted) patient samples, 3 x gt3 and 1x gt1*
  - *four negative controls (EDTA-(mini) pools)*

\* Baylis SA, et al. *J Clin Microbiol.* 2011 Apr;49(4):1234-9.

# HEV NAT quality assessment – Results





# Quality of HEV nucleic acid amplification assays

JOURNAL OF CLINICAL MICROBIOLOGY, Apr. 2011, p. 1234–1239  
0095-1137/11/\$12.00 doi:10.1128/JCM.02578-10  
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Vol. 49, No. 4

## Standardization of Hepatitis E Virus (HEV) Nucleic Acid Amplification Technique-Based Assays: an Initial Study To Evaluate a Panel of HEV Strains and Investigate Laboratory Performance<sup>∇†</sup>

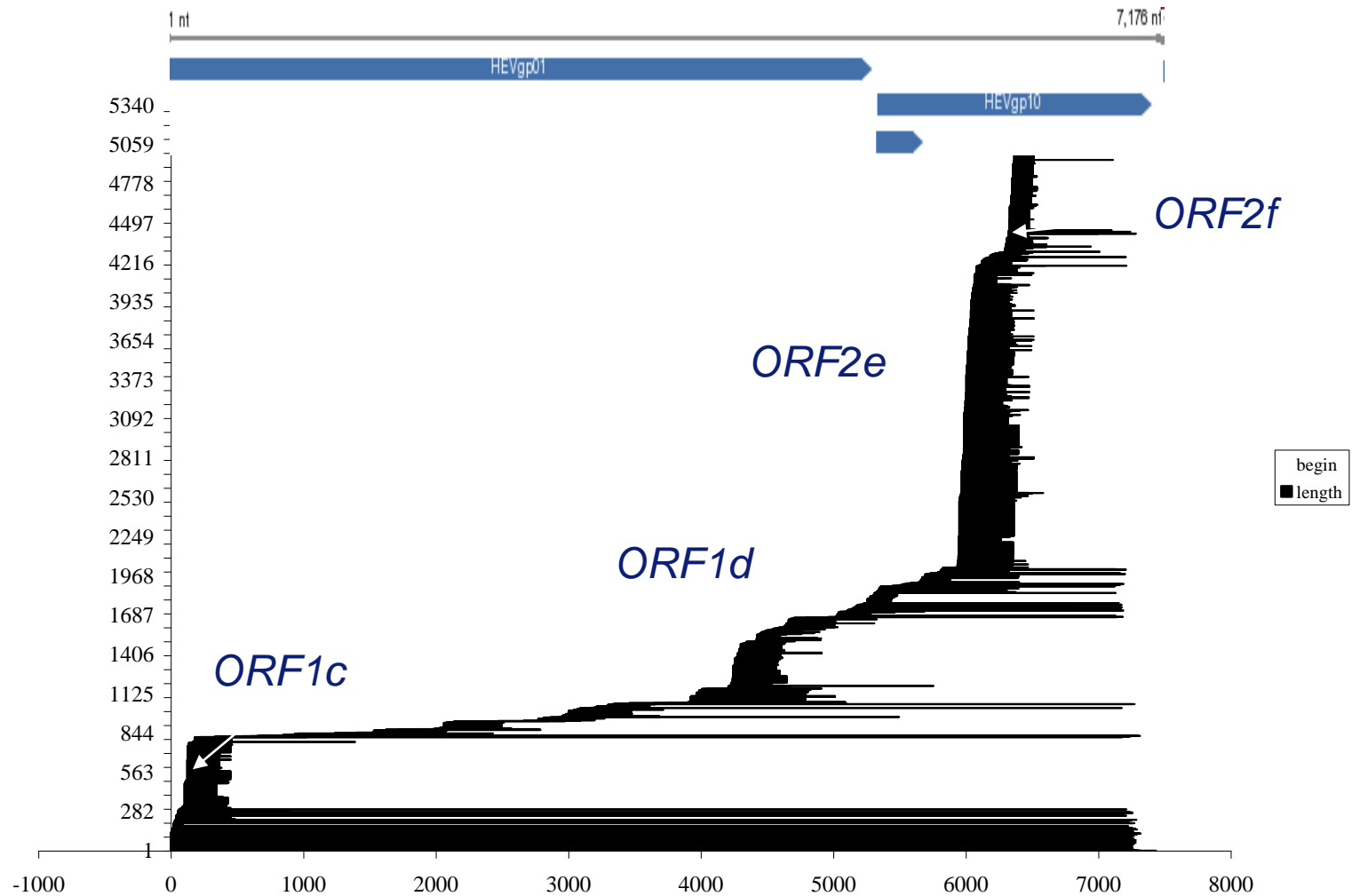
Sally A. Baylis,\* Kay-Martin Hanschmann, Johannes Blümel, and C. Micha Nübling  
on behalf of the HEV Collaborative Study Group‡

24 laboratories, 22 HEV-positive plasma, 10-fold serial dilutions of  
HEV genotypes 3a, 3b, 3f, and 4c.

*International standardisation needed*

→ 1<sup>st</sup> HEV RNA WHO standard available (Gt3, 250,000 IU/ml )!

# Distribution of available sequences along the genome



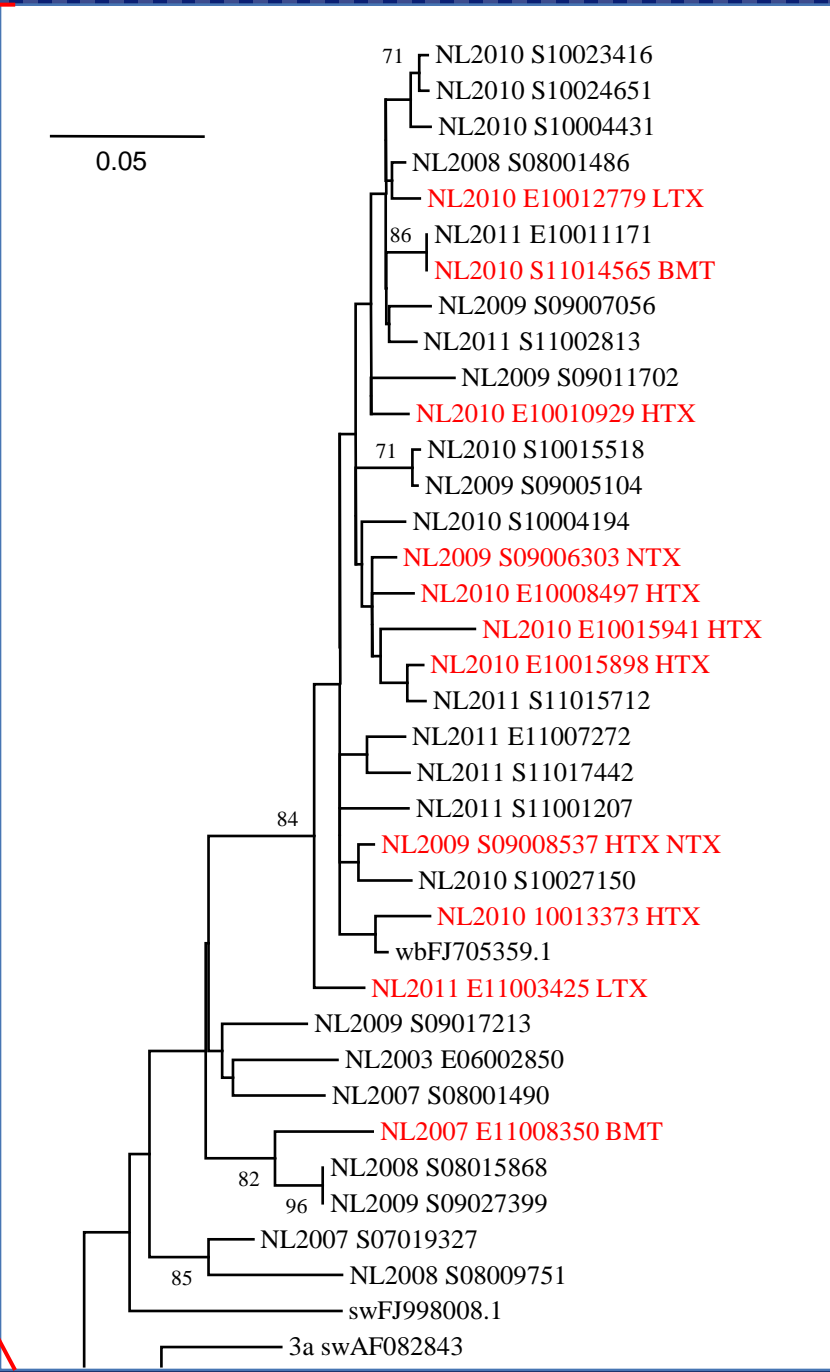
Courtesy: Harry Vennema, RIVM, The Netherlands

SMC

Zafary



ORF1d



asmus MC  
*E. Zafar*

# Conclusies

- Bewustzijn van HEV infecties in transplantatie settings is stijgende, daarom is accurate en snelle diagnostiek nodig voor de juiste interventie strategie
- Om een HEV infectie te diagnosticeren is zowel serologie als moleculaire diagnostiek (real time RT-PCR) van belang
- De nauwkeurigheid van de huidige HEV serologische assays varieert enorm. In onze validatie waren voor IgM DiaPro en Wantai de beste testen, en voor IgG de nieuwe Mikrogen en Wantai assay de beste.
- Kwaliteit van moleculaire assays varieert tussen de verschillende labs, daarom zijn kwaliteitsrondzendingen en standardisatie van belang

# Acknowledgements

## ErasmusMC

*Claudia Mulders*

*Sevgi Deniz*

*Manon Briede*

*Mark Pronk*

*Sandra Scherbeijn*

*Drs. Roel Streefkerk*

*Dr. M. Beersma*

*Prof. Dr. Marion Koopmans*

*Prof Dr. Ab Osterhaus*

*Dr. A. A. van der Eijk*

*Dr. R.A. de Man*

*Dr. P. Th.W van Hal*

*Prof. Dr. W. Weimar*

*Dr. A.H.M.M. Balk*

## RIVM

*Dr. H. Vennema*

*Dr. J. Reimerink*

## Sanquin

*Dr. B. Hogema*