

**Evaluation of the yield of HBV DNA-positive,
seronegative donors using an automated
HIV-1/HCV/HBV triplex NAT assay**

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Ultrio Study Objective

- Evaluate the “yield” of HBV DNA positive donations that are HBsAg and anti-HBc nonreactive using a combination of individual donation and mini-pool (MP) NAT of 16 donations with the Ultrio reagents and TIGRIS platform to supplement the existing Gen-Probe Ultrio license for an HBV DNA screening claim
- Determine rate/characteristics of HBV yield donors
- HBV screening claim approved by FDA on 8/12/08 for MPs up to 16 donations

Scope

- 3 of the 5 ARC NTLs involved (Charlotte, Detroit and St. Louis)
 - All donations tested for HIV-1/HCV/HBV (triplex Ultrio assay) on the TIGRIS instruments using both
 - ID NAT for a target of 600,000 donations
 - MP NAT for a minimum of 2 million donations
 - HIV-1/HCV (duplex) NAT was discontinued during the evaluation
 - Other NTLs continued using HIV-1/HCV (duplex) using the manual platform (eSAS)
 - Testing of WNV assay continued without change
 - TIGRIS or eSAS

Expected Yield

Confirmation of MP NAT Reactivity for ID NAT Yield

- ID NAT: up to 5 window period (WP) donations
 - IND studies: yield of 1:100,000-1:200,000
- MP NAT: 1 additional WP donation
- Dilutional studies to determine reactivity of ID NAT yield samples using pool sizes of: 4, 8, 16

Results (1/28/08-1/5/09)

- Donations tested by MP NAT = 3,118,368
 - 1640 Rx MPs resolved to a Rx donation = **0.05%**
 - Unresolved pool rate = **0.21%**, or 413/194,898
 - Donations tested by ID NAT = 576,490
 - 945 Rx IDs = **0.16%**
 - Total tested = 3,694,858
 - 2585 Ultrio Rx dtns = **0.07%**
 - 2119 (82%) discriminated = **0.06%**
 - 455 nondiscriminated, or **0.01%** (1:8120) of total tested
 - MP NAT = 65 (**1:47,974**)
 - ID NAT = 390 (86%, **1:1478**)
 - 431 eligible for follow up of which 120 have been submitted for reentry
 - 11 QNS for discriminatory testing
-

Discriminated Results (1/28/08-1/5/09)

- 2119 (82%) discriminated, or **0.06%** of total tested
 - 2083 (98%) concordant serologic results from 2060 donors
 - 426 HBV, 231 HIV, 1426 HCV
 - 3 HIV of which **2 confirmed (1:1,847,429)**; 1 false pos
 - 42 HCV of which **15 confirmed (1:246,324)**; 27 false pos
 - 30 HBV of which **9 confirmed (1:410,540)**; 21 false pos
 - 8 MP pos (**1:389,796**), 1 ID pos (**1:576,940**)
 - 6 anti-HBs pos donors with likely vaccine breakthrough (**1:270,956** assuming 44% donors vaccinated; **1:228,680** MP only)
 - 2/5 developed HBsAg; 4/5 developed anti-HBc
 - 3 anti-HBs neg window period donors (**1:689,707** assuming 56% donors unvaccinated; **1:873,143** MP only)
 - 1/2 developed HBsAg; 2/2 developed anti-HBc
-

Results of Dilutions for 019 Yield Donor S/CO values

	<i>Undilute</i>	<i>1:4</i>	<i>1:8</i>	<i>1:16</i>
ARC dHBV TIGRIS	26.55	0.06	0.14	0.19
GP dHBV eSAS	23.28	19.49	0.03	0.03
GP Ultrio eSAS	13.10	0.08	9.81	0.09

HBV Yield Demographic/Risk Info

Donor	Donor Status	Sex	Age	City/State	Risk Factors/Comments
013 MP+, anti-HBs+	Repeat 11/18/05	M	27	Ann Arbor/MI	Donor received HBV vaccine 2000-2001; sexual partner HBV chronic carrier; donor \geq 34 days HBV DNA (100-200 copies/mL); no HBsAg/anti-HBc in >9 months f/u
042 MP+, anti-HBs+	Repeat 2/10/07	M	28	Olmsted Falls/OH	Paramedic in urban setting; donor received HBV vaccination; \geq 75 days HBV DNA (200-45,000 copies/mL); HBsAg SC at day 75 (54 days); anti-HBc SC at day 107
003 MP+, anti-HBs-; strong seroconv	Repeat 5/21/05	F	37	Marietta/GA	HBV vaccine in 1980s; sexual partner HBV chronic carrier; donor 44 days HBV DNA (200-4800 copies/mL); anti-HBc SC at day 70; no HBsAg for 320 days f/u
019 ID+ (MP-), anti-HBs -	Repeat 1/15/08	F	44	White House/TN	Donor denies risk factors and no history of HBV vaccination; refused to enroll in f/u; 100 copies/mL at index
011 ID+ (MP+), anti-HBs+	Repeat 3/29/07	F	17	Centralia/IL	Donor received HBV vaccine 10-11 yrs prior to index donation; sexual partner HBV chronic carrier; donor \geq 137 days HBV DNA (100-50,000 copies/mL); HBsAg SC at day 108 (60 days rx); anti-HBc SC at day 168

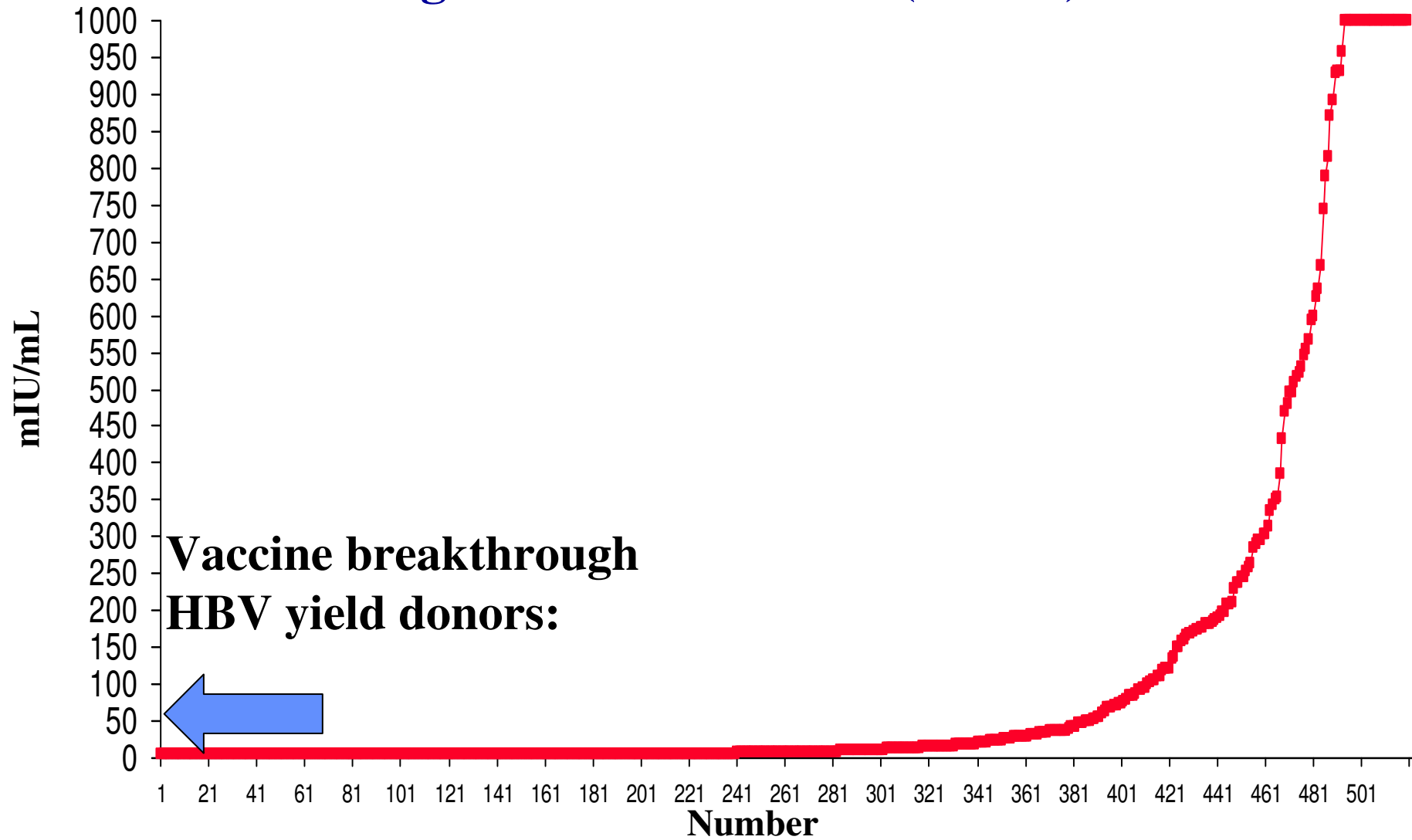
HBV Yield Demographic/Risk Info

Donor	Donor Status	Sex	Age	City/State	Risk Factors/Comments
055 MP+, anti-HBs-	First-Time	M	20	Batesville/MS	Donor in Job Corps; no history of HBV vaccine (vaccinated after finding out of DNA+ result); ≥ 75 days HBV DNA with high-titer viremia ($200-10^8$ copies/mL); HBsAg SC at day 41 (123 days rx); anti-HBc SC at day 70
074 MP+, anti-HBs-	Repeat 05/17/08	F	24	Coral Springs/FL	Iranian donor; cannot recall HBV vaccination history; sexual partner HBV chronic carrier; HBV DNA ≥ 73 days (200 copies/mL); no HBsAg for >116 days f/u; anti-HBc SC at day 73
001 MP+, anti-HBs+	Repeat 03/31/08	M	22	Boston/MA (NY)	College student who received HBV vaccine 1991-1992; HBV DNA ≥ 49 days (viral load pending); sexual partner HBV chronic carrier no HBsAg for >91 days f/u; anti-HBc SC at day 69
029 MP+, anti-HBs+	Repeat 04/03/07	M	19	Colerain/NC	HIV infected donor (RNA + Ab); unable to be contacted for participation in f/u and confirmation of vaccination

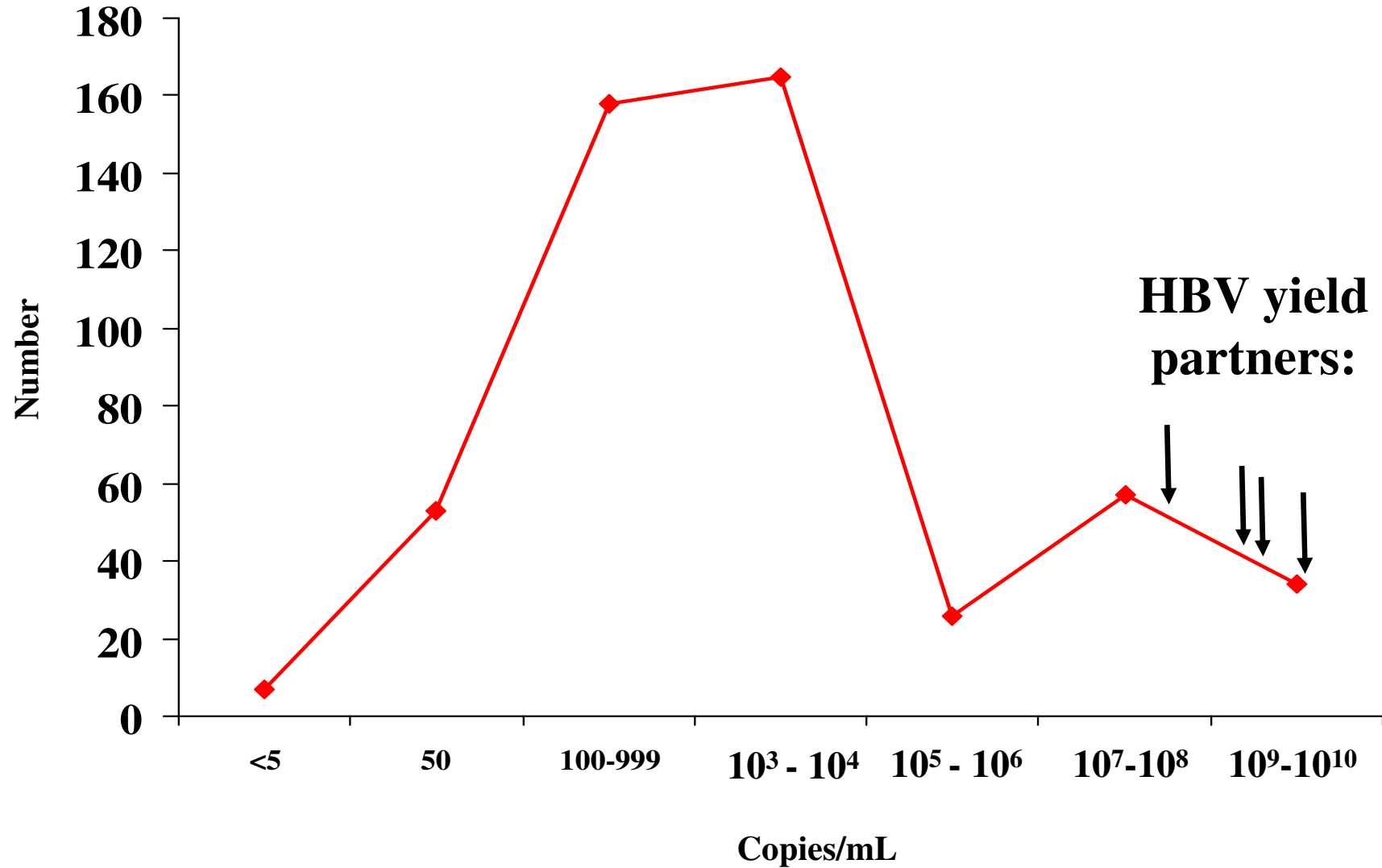
Virologic/Serologic Markers in HBV DNA Yield Donors

Donor	Anti-HBs mIU/mL @ index	Time (days) followed	Viral Load Range (c/mL)	Duration (days) DNA pos	HBsAg first pos day (duration)	Anti- HBc first pos day	Anti- HBc IgM
013	+; 43	243	100-200	≥ 34	-	-	-
042	+; 33	133	800-45,000	≥ 75	75 (54)	107	+
003	+; 3	320	200-4800	= 44	-	70	+
011	+; 11	189	100-50,000	≥ 137	108 (60)	168	+
055	-	>175	>10 ⁸	≥ 75	41 (123)	70	+
074	-	>116	200	≥ 73	-	73	+
001	+; 100 (day 45)	>115	100	≥ 49	-	69	+

Anti-HBs Concentrations in HBsAg and Anti-HBc Negative Blood Donors (N=520)



HBV DNA Viral Loads among HBsAg and anti-HBc Positive Donors (N=500)



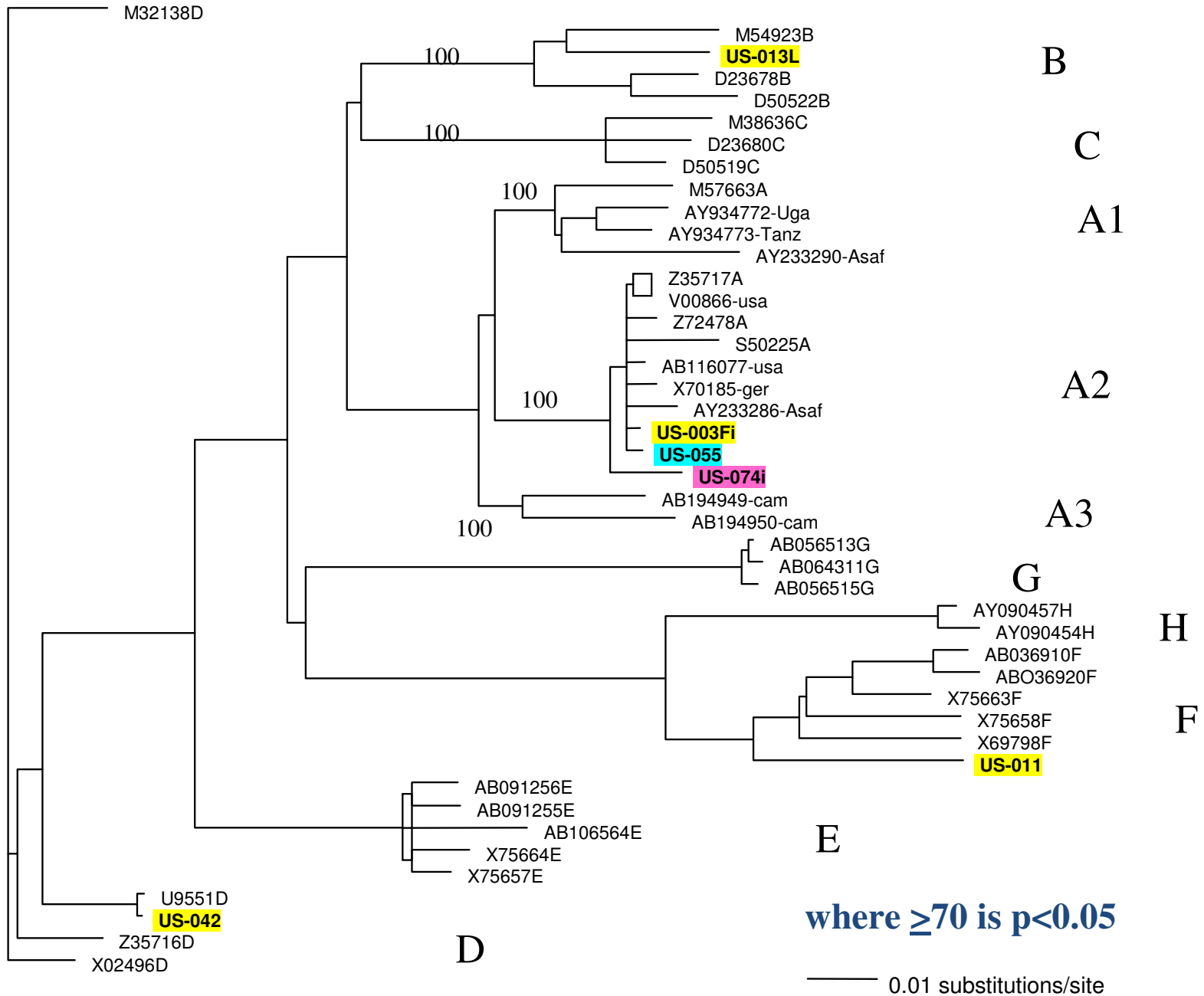
HBV Additional Studies

- Performed by 3 groups
 - Prof Wolfram Gerlich
 - Institute for Medical Virology, Univ of Giessen, Germany
 - Viral loads, genotype, subtype, sequence analysis
 - Dr. Paul Coleman
 - Abbott Laboratories
 - Anti-HBc IgM (ARCHITECT), sequence analysis
 - Prof JP Allain
 - Laboratory of Molecular Virology, Depart Haematology, Univ of Cambridge, UK
 - Viral loads, genotype, sequence analysis
 - Sequencing
 - Full length, preS/S, BPC/PC

Sequencing of HBsAg with Genotype and Subtype

Donor or Partner	DNA Viral Load (copies/mL)	HBsAg Conc. (ng/mL)	Genotype	Subtype	Mutation (No. and Location)
Donor - 003	1,200, 86		WT A2 (Central Europe, N Amer)	<i>Pending</i>	5 - sN59D; sT143A; sS210N; sL22S; sK122E
Partner - 003	2.7 x10 ⁹	67,300	WT A2	<i>Pending</i>	2 – preS V172M and rtC6Y
Donor - 011	2,700, 13		WT F1 (American Indians of Central America)	<i>Pending</i>	2 - S and preS domain stop codons @ sY71 Stop and preS F24L
Partner - 011	2.4 x10 ¹⁰ , 2.6 x 10 ⁸	100,600	WT F1	<i>Pending</i>	1 - T173M Not present in other F1 strains
Donor - 013	35		WT B2 (East Asia)	adw2	1 - sF220L (rtL229V) Not present in other B2 strains
Partner - 013	8.0 x10 ⁹ , 1.8 x 10 ⁶		WT B2	adw2	1 - sF220L (rtL229V) Not present in other B2 strains
Donor - 042	65, 230, 43		WT D3 (Mediterranean) (preS domain)	ayw3	2 - sT125M and sP127T
			WT A2 (S domain)	ayw3	2 - sG44R and sT143A
Donor - 001	11		C2 (N China, Korea)	adw2	1 – G145R; vaccine escape mutation
Partner - 001	1.9 x 10 ⁸		C2)	adw2	1 – G145R; vaccine escape mutation

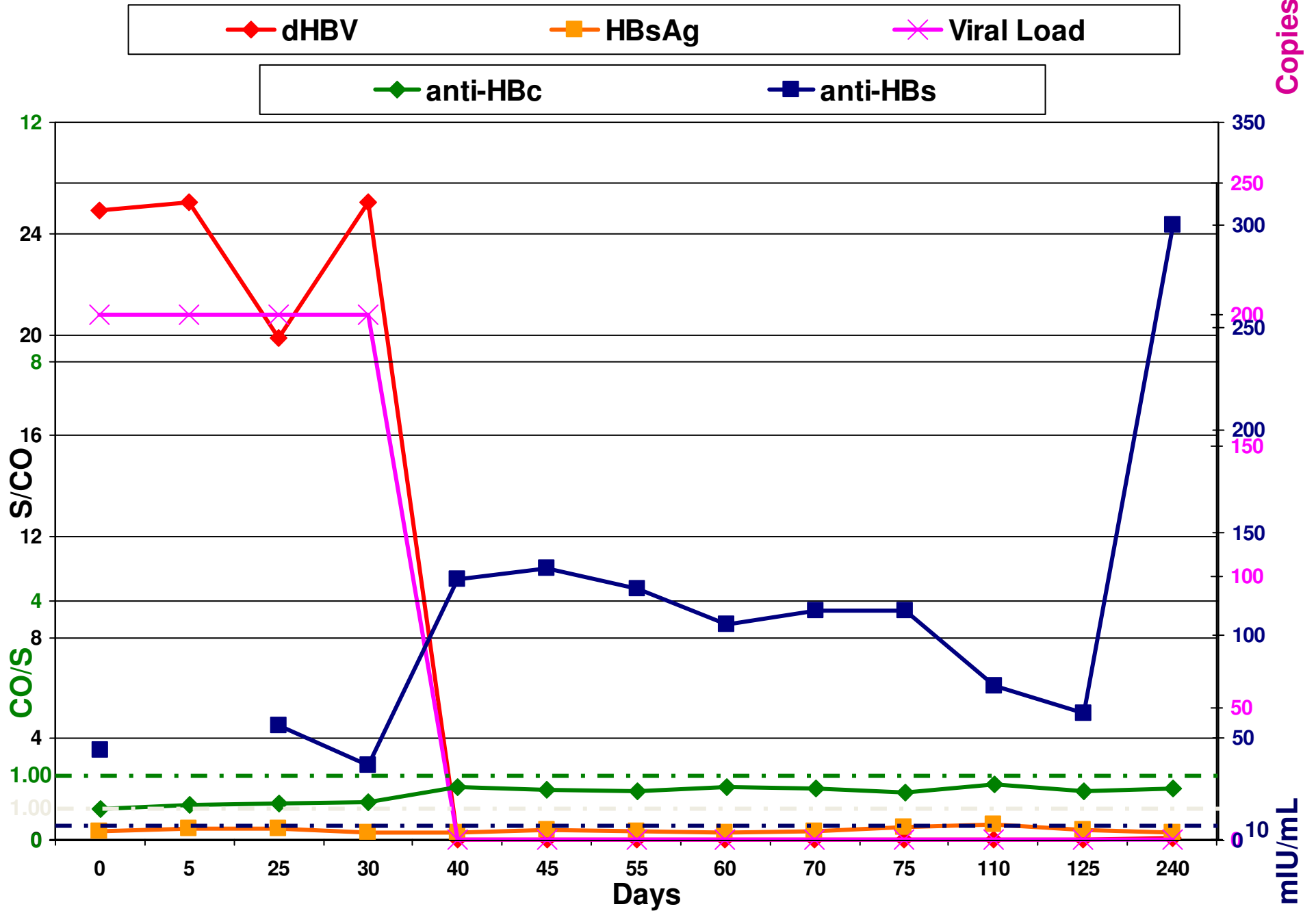
HBV Bootstrap Analysis for 6 cases with full genome sequences



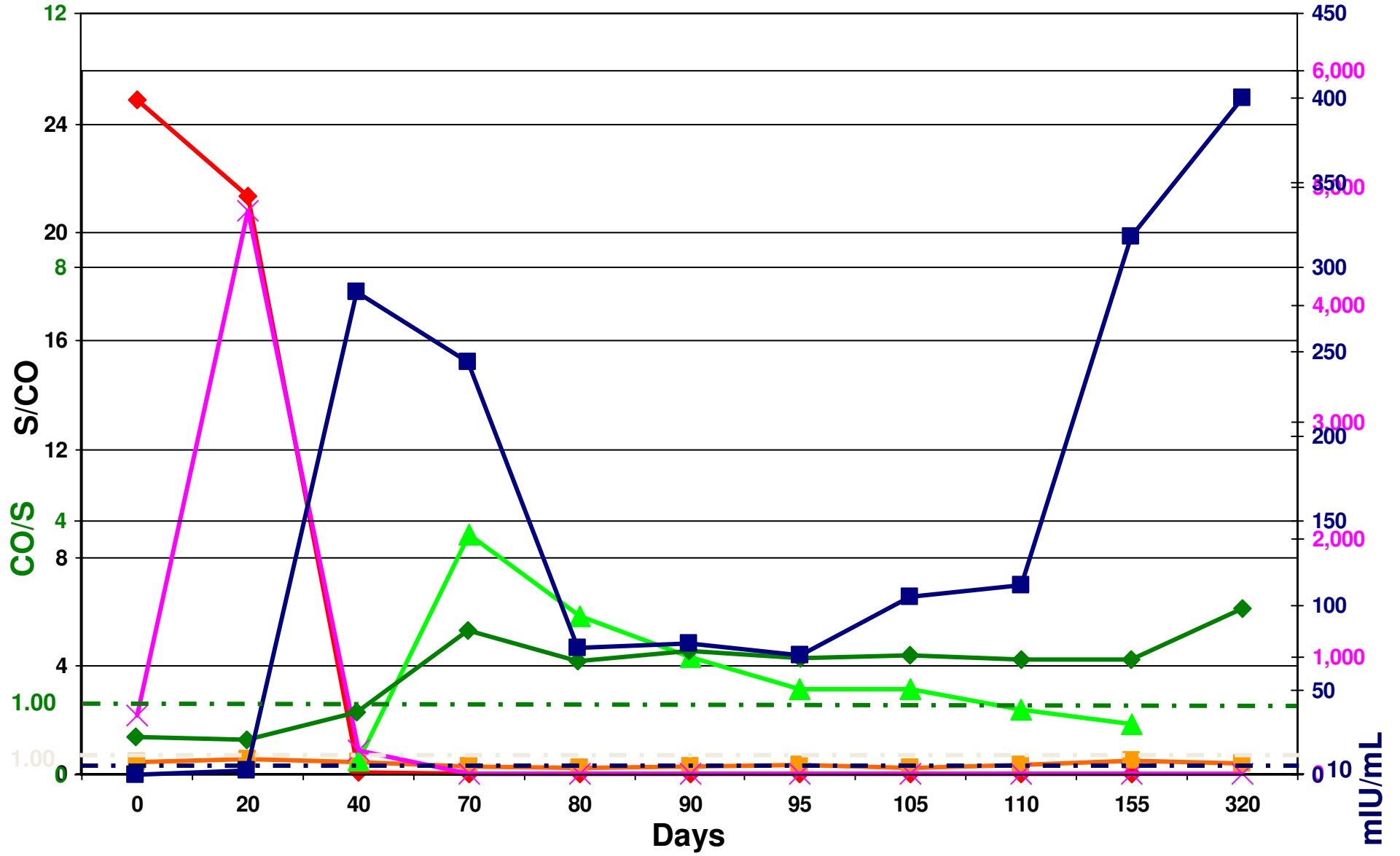
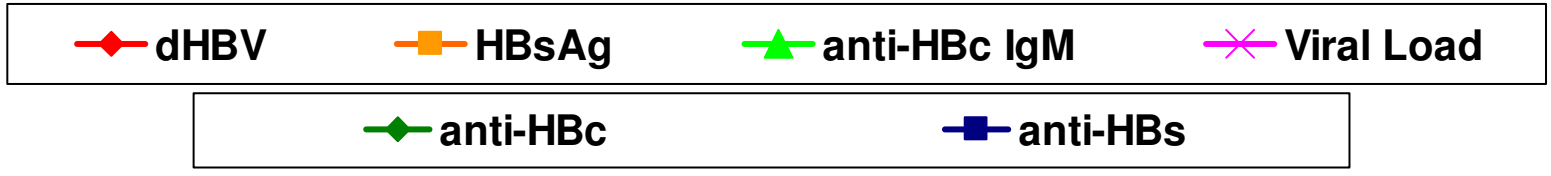
Virologic/Serologic Profiles

- Breakthrough cases with likely infection via a sexual partner
 - All partners tested and confirmed to have high levels of HBV DNA ($>10^7$ copies/mL), HBsAg, anti-HBc but no anti-HBs, and when sequencing complete, same virus as the donor
 - Donors with and without HBsAg development
- Breakthrough case occupational exposure
- Acute infection in the absence of vaccination

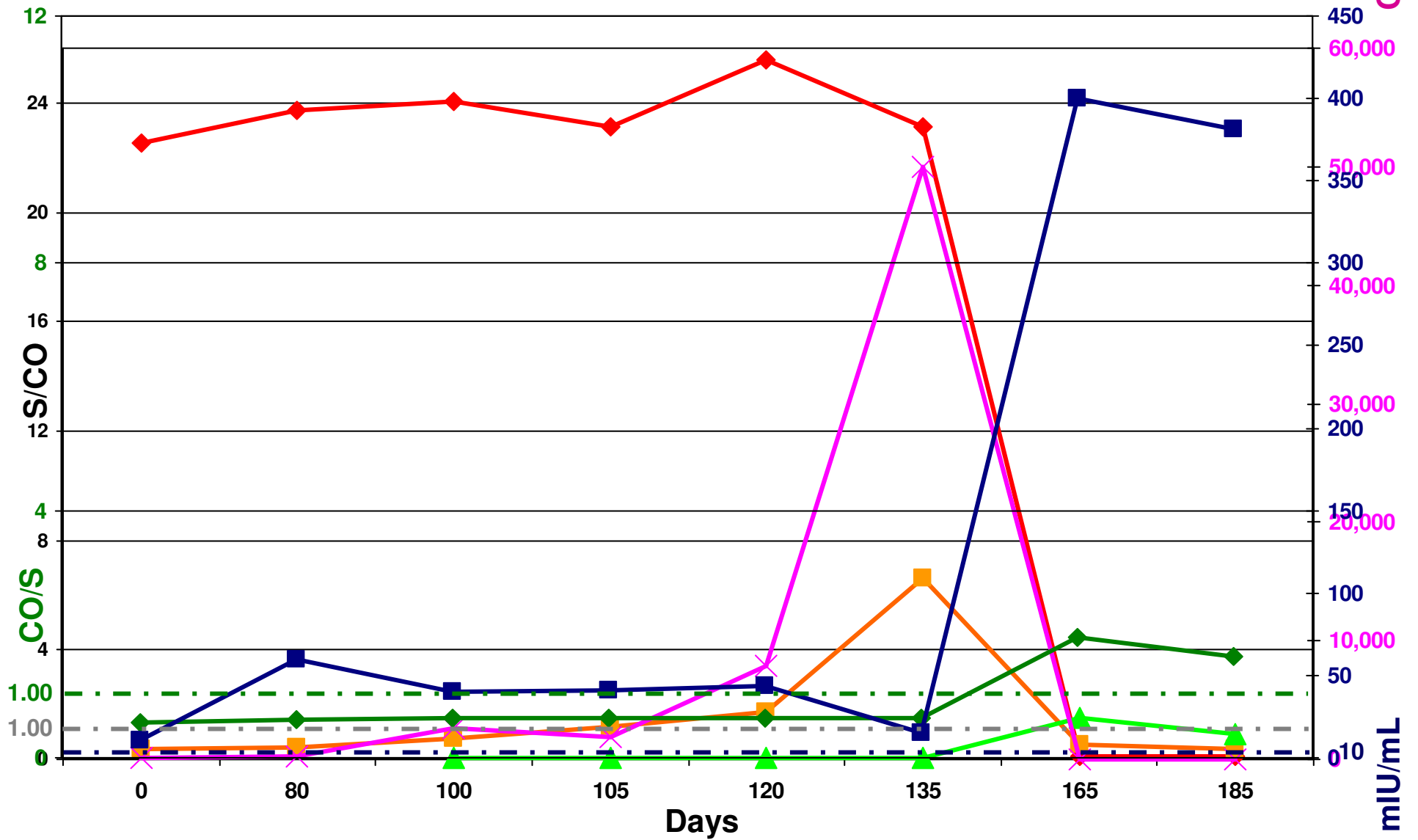
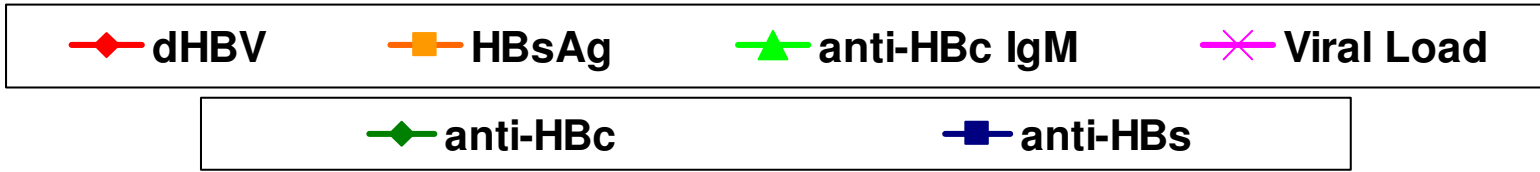
UFS Yield Donor 013KF17011



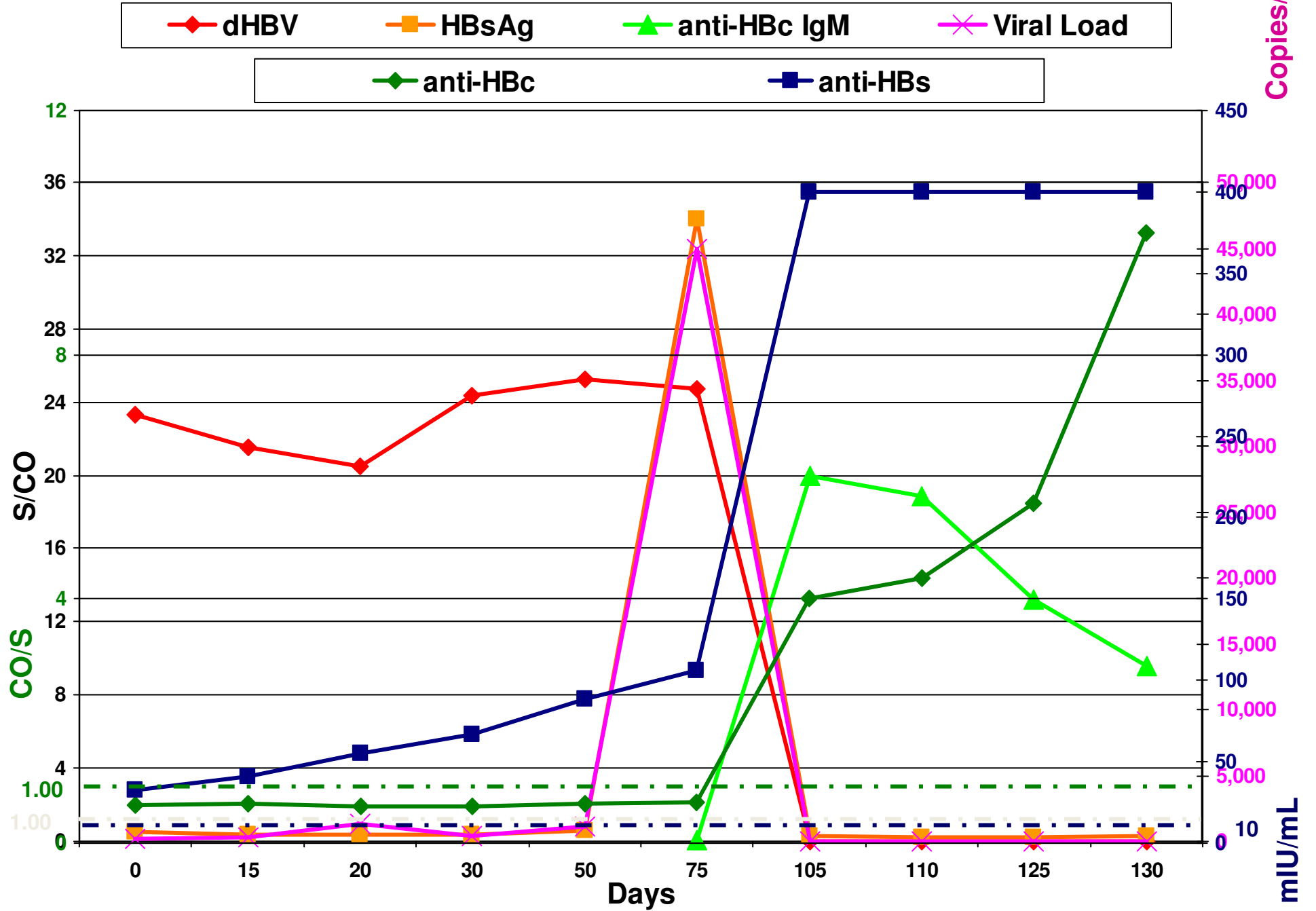
UFS Yield Donor 003FQ75130



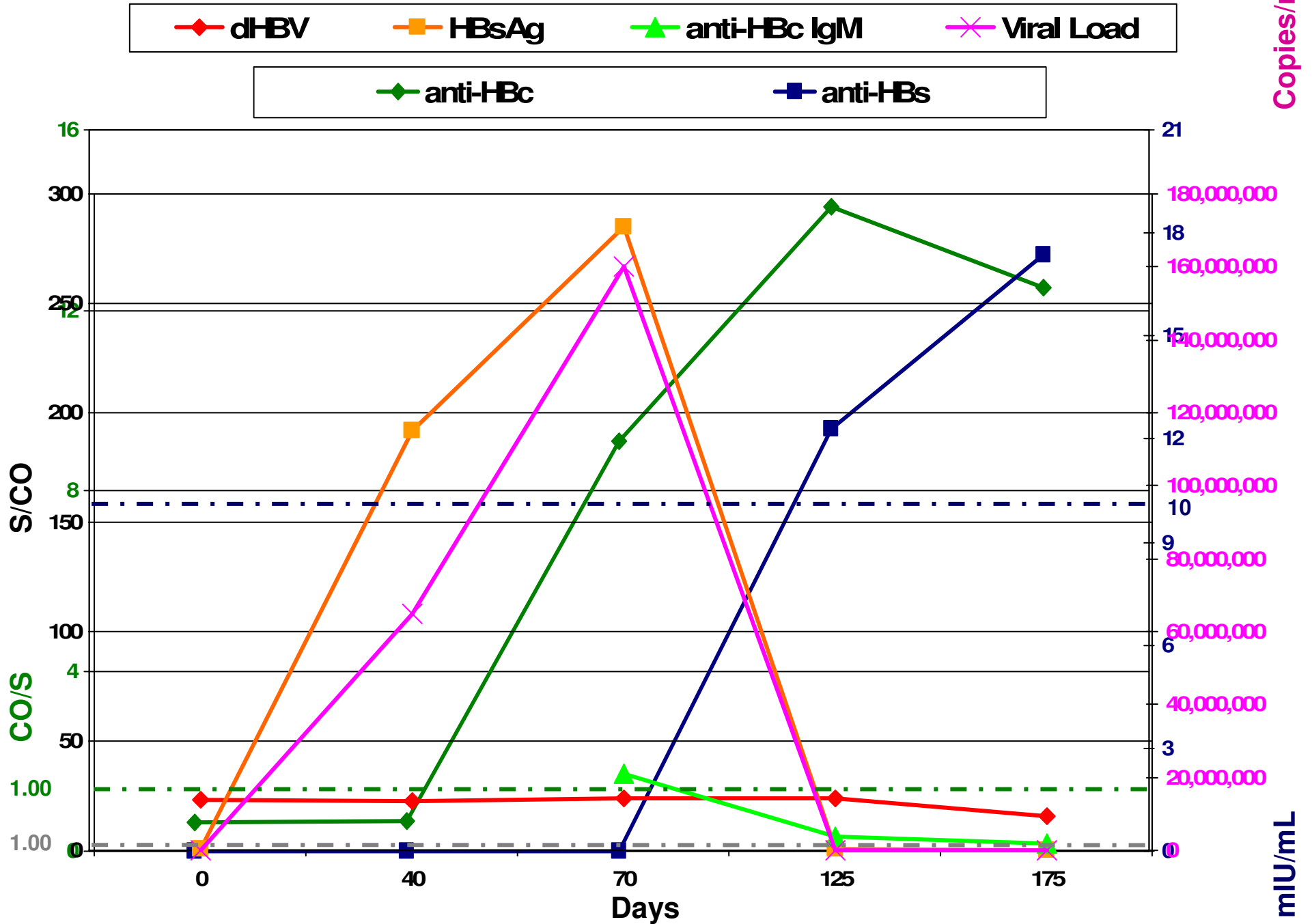
UFS Yield Donor 011KC34573



UFS Yield Donor 042FM54241



UFS Yield Donor 055N 30971



Comparison of HBV DNA Yields in US Studies

ABC Roche Trials (1, 6, 24)

1:352,451

(2/704,902) COBAS AmpliScreen trial

1:425,730

(7/2,980,103) + extended COBAS AmpliScreen testing

1:72,336

(1/72,336) MPX trial

1:381,555

(8/3,052,439) **Combined**

1:610,488

(5/3,052,439) **Combined** (*- PRISM HBsAg RRs*)

ABC GP Trial (1, 8) 9 sites; 5/07 to 7/16/08

1:282,984

(4/1,131,937) vs GSC 3.0 and PRISM HBsAg

ARC GP Trial (1, 16) 1/29/08 to 1/5/09

1:410,540

(9/3,694,858) vs PRISM HBsAg

Summary

- Performance of duplex (eSAS) and triplex (ULTRIO[®]-TIGRIS) is comparable regarding MP NAT specificity and HIV-1/HCV NAT yield
 - 9 HBV DNA pos/HBsAg and anti-HBc neg donations produced comparable yields to other HBV yield studies performed in the US
 - **1:410,540** yield rate
 - 8 of 9 yield donors were detected by MP NAT; **1:389,796** yield rate
 - 7 with long-term follow up
 - All with detectable HBV DNA for 34-137 days
 - 3 with HBsAg at 41-75 days after DNA (duration 54-123 days)
 - 6 with anti-HBc SC at days 69-168
 - 6 of 9 were immunized individuals having anti-HBs at index or shortly thereafter; although these represent acute infections, dynamics of infection differ, and infectivity of such donations is unknown; **1:270,956** yield rate (or **1:228,680** for MP only)
 - 3 of 9 were window period donors; **1:689,707** yield rate (or **1:873,143** for MP only)
-

Conclusions re HBV NAT

- ID NAT modeling studies (nonvaccinated donors)
 - Lowest residual risk; highest yield = **1:466,000 – 1:713,000**
 - Logistically not feasible for either platform
 - Modeling cannot predict total yield due to vaccinated donors with differing kinetics early in infection
- MP NAT modeling studies (nonvaccinated donors)
 - MPX: MP 6 highest yield = **1:830,000 (= 1:713,000)**
 - Ultrio: MP 8 = MP 16 = **1:1,345,000 – 1:2,000,000**
- MP NAT observed yield (both vaccinated and nonvaccinated)
 - MP NAT at various pool sizes = **1:300,000-1:600,000**
 - Data do not suggest benefit of MP 8 vs MP 16
 - ARC study; MP 16 detected 8 of 9 units
 - 9th yield unit **only** detected by ID NAT (not detected MPs 4, 8 or 16)
- High yield of MP 16 indicates value in comparison to ultra-sensitive HBsAg assays
 - Total yield = **1:410,540**
 - MP 16 yield = **1:389,796**
 - Vaccinated MP 16 yield = **1:228,680**; infectivity ?
 - Nonvaccinated MP 16 yield = **1:873,143**

**Are anti-HBs positive units
infectious?**

HBV Transmission from Low Level HBV DNA Pos “Occult” Donor

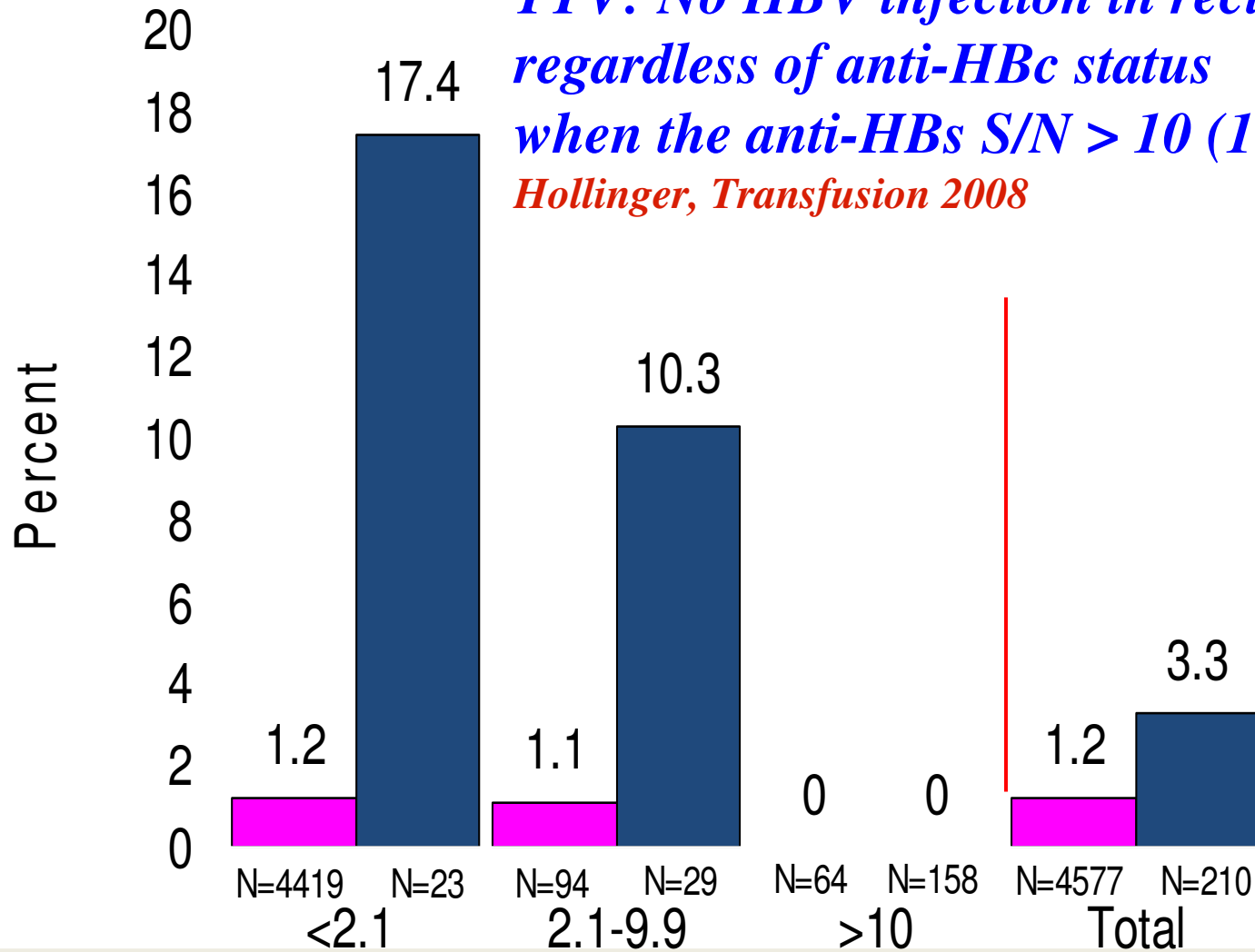
Levicnik-Stexinar et al., J Hepat 2008:48

	HBsAg	Anti-HBc	Anti-HBc IgM	Anti-HBs (mIU/mL)	HBV DNA (IU/mL)	Genotype
Donor (index)	Neg	Pos	NT	12-29	180	D
Recip* #1 4 days	Pos	Pos	Pos	Neg	NT	
8 days	Neg	Pos	Pos	Neg	185	D
Recip #2 7 days	Pos	Neg	Neg	Neg	1.1x10⁸	D
14 days	Neg	Pos	Pos	Neg	Neg	

* Neg pretransfusion

■ Anti-HBc Neg ■ Anti-HBc Pos

TTV: No HBV infection in recipients regardless of anti-HBc status when the anti-HBs S/N > 10 (10-60 mIU/mL)
Hollinger, Transfusion 2008



Donor Anti-HBs S/N Value



Results of JRC LB Study

Component Infectivity from ID NAT + Repository					
Anti-HBc in donors	No. donors	No. products tested	Infection		Infection Rate
			yes	no	
Low titer	29	33	1	32	3%
Negative	20	22	11	11	50%
Unknown	6	8	0	8	
Total	55	63	12 (100-380 c/mL)	51	19%

**Of 12 infectious components, 11 (92%) were window period
and 1 (8%) was anti-HBc positive (jumbo FFP)**

Results of JRC LB Study

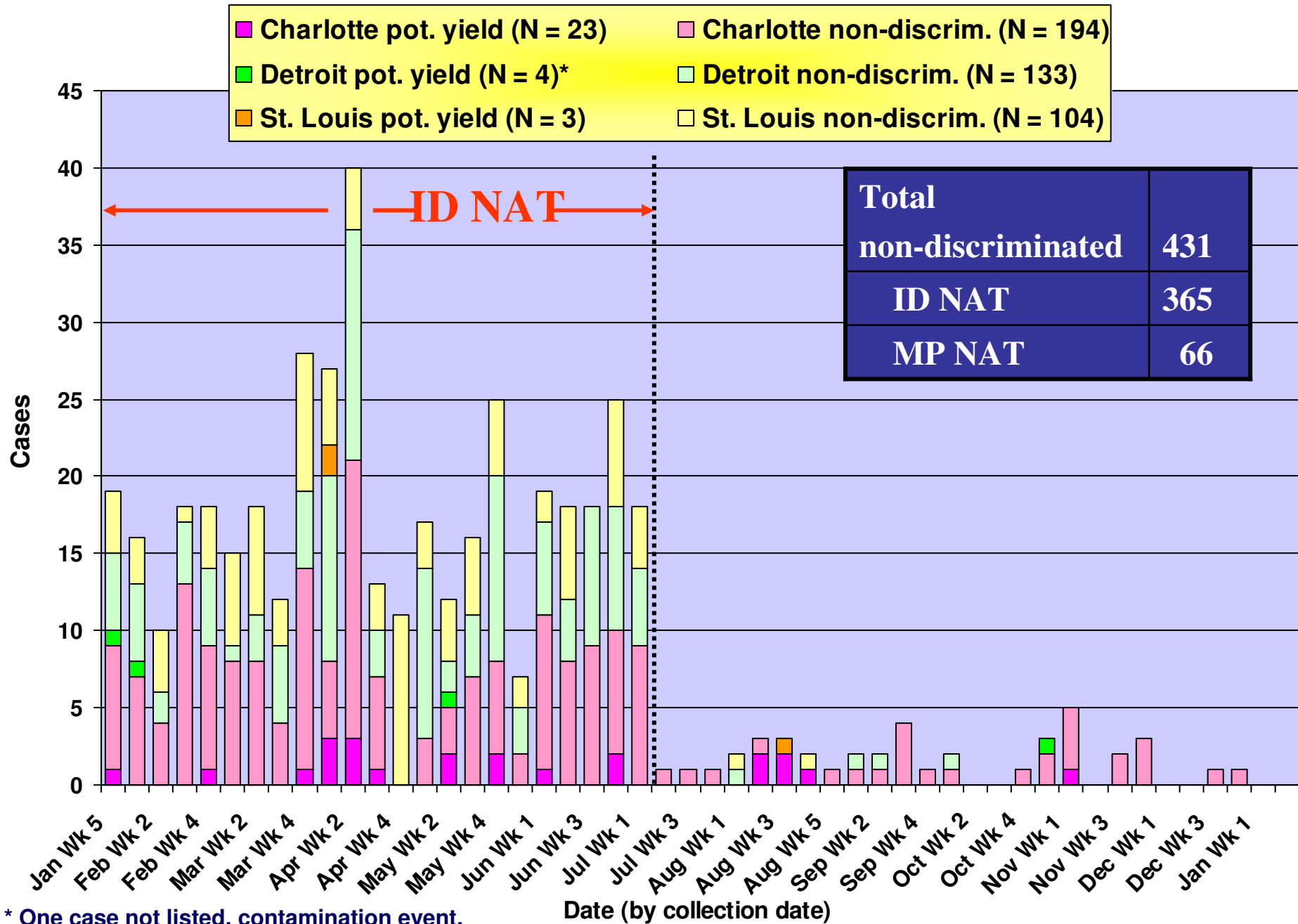
Anti-HBc in donors	Infectious	Anti-HBs in donors	
		Pos	Neg
Low titer	Yes	0	1
	No	11	12
Negative	Yes	0	8
	No	0	8
Unknown	Yes	0	0
	No	5	2
Total	Yes (12)	0	9 (29%)
	No (51)	16 (100%)	22

Additional Slides for questions

Additional Results (JPA) and Conclusions to Date

- Full sequences obtained for 6 (013, 003, 055, 074, 011, 042)
 - 1 each F, B, D and 3 each A2; half representing genomes not typically found in the US (majority in US is A2 in Caucasians, A1 in Blacks and B/C in Asians)
- Pre-S/S or BCP/PC sequences for 8 (above + 019 + 001)
 - 7 WT (compared to ref strains in GenBank); all major hydrophilic regions (MHRs) of the S protein conserved. Also, core, pol and X genes conserved; no stop codons
 - 1 S protein vaccine escape mutation (001)
- 4 (013, 003, 011, 042) are likely breakthrough infections, since no MHR mutations and all WT; one (001) vaccine escape
 - Inability of background levels of vaccine-derived anti-HBs to neutralize infectious virus
 - Low level anti-HBs might be sufficient to complex HBsAg making it undetectable but not sufficient enough, or too low, to prevent infection and the detection of HBV DNA
 - Public health implications; vaccine does prevent chronic HBV but not infection (all of these donors did clear virus and had truncated HBsAg responses, if present); more frequent boosters may be warranted
- 3 likely window period cases (019, 055 and 074)

Impact of ID NAT



* One case not listed, contamination event.

Specificity Comparisons

Duplex eSAS – MP NAT (16); 3 NTLs to 12/07

➤ 33,868,522 screened = 99.9974% (99.9972%-99.9975%)*

Ultrio TIGRIS – MP NAT (16); 3 NTLs

➤ 3,118,368 screened = 99.9973% (99.9966%-99.9978%)*

Ultrio TIGRIS – ID NAT; 3 NTLs

➤ 576,490 screened = 99.9297% (99.9225%-99.9364%)*

*95% CIs by the binomial distribution

Concordant Ultrio/Serology

2083 Results/2060 Donors

No. Reactive	HBV	HIV	HCV (+anti-HBc)
St Louis	63*	23	333*
Detroit	78	22*	222*
Charlotte	285* (67%)	186* (81%)	871* (61%)
Total	426	231	1,426**
Rate	1:8673	1:15,995	1:2591

*12 donations reactive for both HIV and HCV Ab + NAT
 5 donations reactive for both HBV and HCV Ab + NAT
 6 donations reactive for both HBV and HIV Ab + NAT

** 327 (23%) HCV reactives also anti-HBc reactive



Ultrio Nonreactive – Serology Reactive (Confirmed Positive); N=5666 from 5662 Donors

	HBsAg (+)	Anti-HCV (+)	Anti-HIV (+)
ID NAT (-) (N=576,490)	134 (25)	455 (71)	339 (2)
MP NAT (-) (N=3,118,368)	655 (153)	2615 (446)	1468 (27)
TOTAL	789 (178)	3070 (517)	1807 (29)
Rate False Pos Serology (ID or MP NAT -)	1:5289 1:6212	1:1501 1:1438	1:1710 1:2164
Rate of Conf'd Pos Serology (ID or MP NAT -)	1:23,060- 1:20,381	1:8120 1:6992	1:288,245- 1:115,495

Comparison of HBV DNA Reactivity in Anti-HBc Reactive Donors

(with removal of all HBsAg, HCV and HIV reactives)

	22 Reactive Ultrio MP NAT of 6573 anti-HBc RRs	12 Reactive Ultrio ID NAT of 1447 anti-HBc RRs	34 Total Reactive of 8020 anti-HBc RRs
No. (%) dHBV Pos	18 (0.33%)	9 (0.62%)	27 (0.34%)
(No. PCR* Pos/No. Tested)	(5/9 = 56%)	(6/7 = 86%)	(11/16 = 69%)
Adjusted	10 (0.15%)	8 (0.55%)	19 (0.23%)

*Multiprep Ampliscreen HBV ID NAT



HBV Sensitivity of NAT Systems according to the PIs

- ID NAT Ultrio/TIGRIS = 10.40 IU/mL
- MP NAT 8 Ultrio/TIGRIS = 83.20 IU/mL
- MP NAT 16 Ultrio/TIGRIS = 166.40 IU/mL

- ID NAT MPX/s 201 = 3.80 IU/mL
- MP NAT 6 MPX/s 201 = 22.80 IU/mL

**Table 1 HBV DNA Yield Samples
Index and Follow up Results
Linauts et al. Transfusion July 2008**

	DRAW DATE	INDEX SAMPLE – HBsAg & anti-HBc NR		FOLLOW UP SAMPLE: DAYS REACTIVE (DAYS OF FOLLOW UP)					DONOR HISTORY
		anti-HBs mIU/mL	HBV DNA c/mL	HBsAg	anti-HBc IgM	anti-HBc Total	anti-HBs	DNA	
1	6/11/2003	51	61,000	7 (54)	26-54 (54)	26-54 (54)	0-54 (54)	0-7 (54)	27 YEAR OLD MALE REPEAT DONOR-MULTIPLE MALE SEXUAL PARTNERS
2	4/24/2004	NT	37,000	14 (177)	NR to 177	28-177 (177)	<14- 177 (177)	0-21 (177)	50 YEAR OLD MALE REPEAT DONOR, NO RISK FACTORS IDENTIFIED
3	7/24/2003	NT	2300	7-21 (134)	28-63 (63)	28-134 (134)	63-91 (91)	0-28 (134)	29 YEAR OLD MALE REPEAT DONOR ACUPUNCTURE 8 WEEKS PRIOR TO DONATION
4	9/6/2002	NT	2000	17-200 (200)	55-200 (200)	48-200 (200)	NR to 200	0-200 (200)	26 YEAR OLD MALE REPEAT DONOR, NO KNOWN RISK



**Table 1 HBV DNA Yield Samples
Index and Follow up Results
Linauts et al. Transfusion July 2008**

	DRAW DATE	INDEX SAMPLE – HBsAg & anti-HBc NR		FOLLOW UP SAMPLE: DAYS REACTIVE (DAYS OF FOLLOW UP)					DONOR HISTORY
		anti-HBs mIU/mL	HBV DNA c/mL	HBs Ag	anti-HBc IgM	anti-HBc Total	anti-HBs	DNA	
5	10/24/2006	NR	270						UNABLE TO ENROLL IN FOLLOW UP
6	6/11/2005	NR	200						39 YEAR OLD MALE REPEAT DONOR REPORTED VACCINATION HISTORY 3 DAYS PRIOR TO DONATION. COULD NOT CONTACT FOR F/U;
7	9/25/2002	2340	200	NR to 167	29-167 (167)	22-167 (167)	0-167 (167)	0-22 (167)	49 YEAR OLD FEMALE REPEAT DONOR, HISTORY OF VACCINE BUT WITH NEGATIVE TITER 8 WEEKS PRIOR TO INDEX
8 MPX*	7/10/2005	RR	<LOD	NR to 59	NT	NR to 59	0 (0)	0-59 (59)	PHILLIPPINE IMMIGRANT; NO HISTORY OF VACCINATION

* Nonreactive by MP COBAS AMPLISCREEN; Reactive IDT using multiprep procedure



American Red Cross

***Table 2 Window Period Samples
Repeat Viral Load, Genotype and PRISM Results
Linauts et al. Transfusion July 2008***

Donor #	Draw Date	HBV DNA Quantitation original copies/mL	HBV DNA Quantitation 2007 repeat copies/mL	HBV genotype	PRISM S/CO
1	6/11/2003	61,000	31,000	H	1.22, 1.16, 1.16
2	4/24/2004	37,000	3850	C	0.63
3	7/24/2003	2300	2480	B	1.90, 1.99, 1.97
4	9/6/2002	2000	ND	ND	1.11, 1.23, 1.09
5	10/24/2006	270	<200	NA	0.22
6	6/11/2005	200	<200	NA	0.81
7	9/25/2002	200	ND	NA	0.22
8	7/10/2005	<LOD	<200	NA	0.19