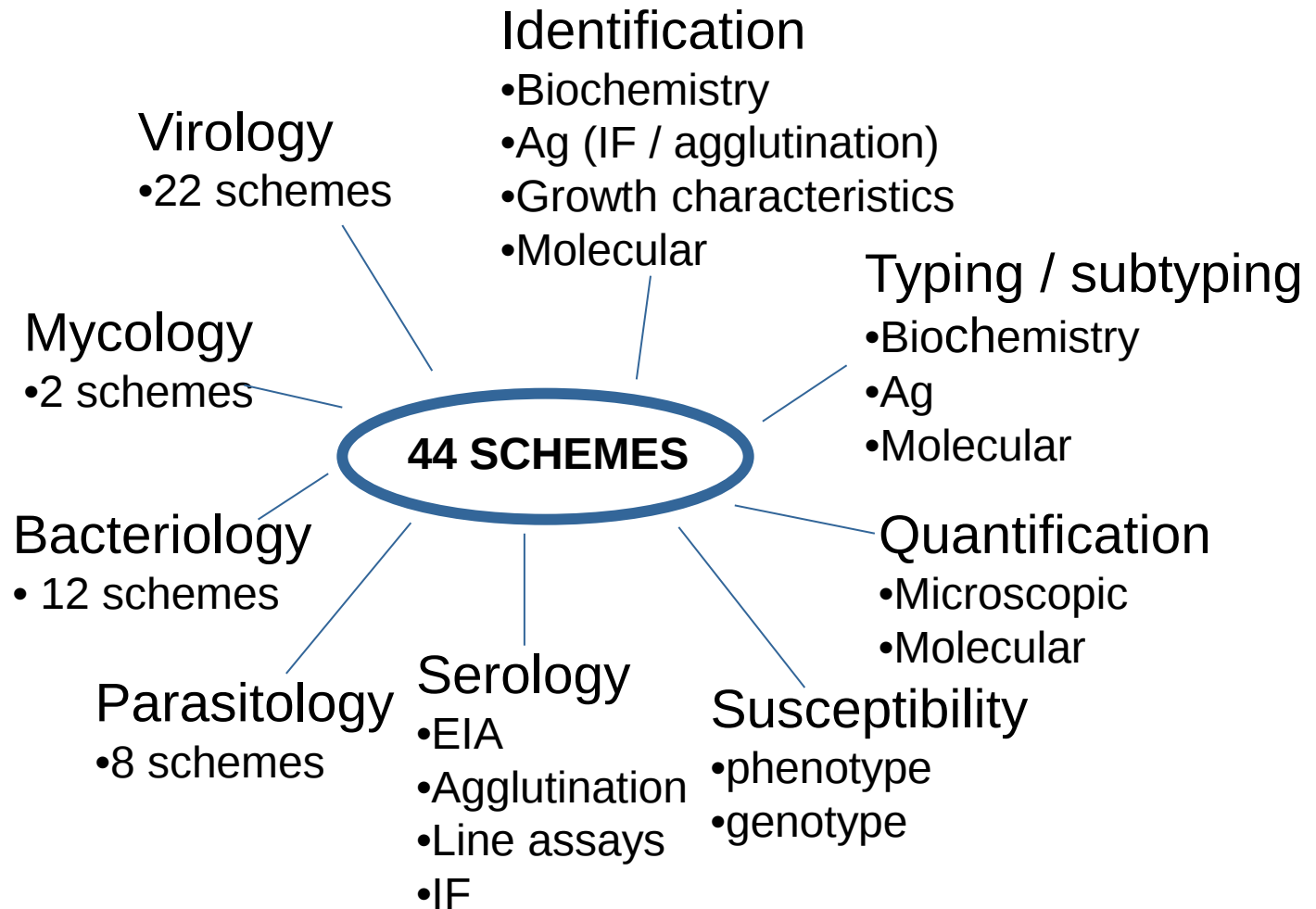
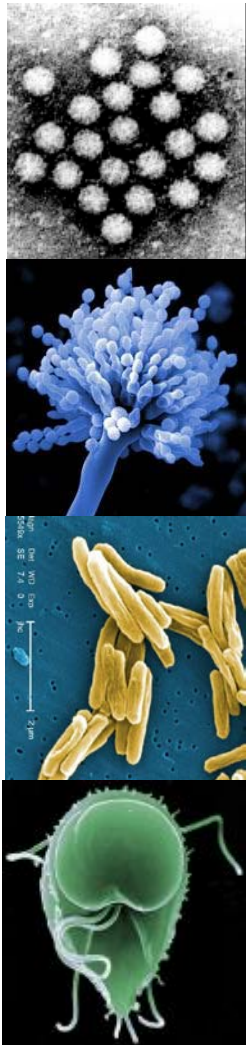


Impact of the introduction of the International Standards: evidence from EQA panels

Dr Beatrix Kele
UK NEQAS for Microbiology
NCI Workshop on Harmonization of EBV Testing
for Nasopharyngeal Cancer
5-6 November, 2015

Schemes overview



Introduction: EBV

- EBV infection
- Seroconversion: 90% at young adult
100% by 40 years of age
- Transmission: body fluids, transplantation
- B-lymphocytes are the site of latency

Introduction: Clinical significance

- Primary infection
- Infectious mononucleosis
- Chronic active EBV
- Associated with: Burkitt lymphoma and nasopharyngeal carcinoma (>95%), Hodgkin's disease (<5-70%), gastric adenocarcinomas (2-16%), T/NK cell lymphomas (40->90%)
- Immunocompromised patients
- Clinical diagnosis: **EBV DNA detection**

QUESTIONNAIRE / April 2009

- Sent to CMV DNA quantification scheme participants (N=81)
- 78.8% use molecular methods to quantify EBV DNA (all real time PCR)
- Sample tested:
 - 62.5% use whole blood
 - 54.2% use plasma
 - 16.6% use both
- Median detection limit: 250 copies/mL (10-2,000)
- Median number of samples examined per week: 22.5 (0.5-200)
- Patient types:
 - 68.8% - HSCT
 - 36.8% - SOT
 - 7.9% - tumour
- Clinical cut-off: 'viral load rise over a definite period of time' & 'confirm viral load above the cut-off'

Pilot /Specimen design

	Namalwa/mL	Expected VL		Pre-distribution Artus		Homogeneity
		c/mL	Log	c/mL	Log	SD
A	78,000	156,000	5.19	102300	5.01	0.118
B	7,800	15,600	4.19	31600	4.50	0.149
		Log diff.	1.00	Log diff.	0.49	
				(Qiagen column)		(20 vials)

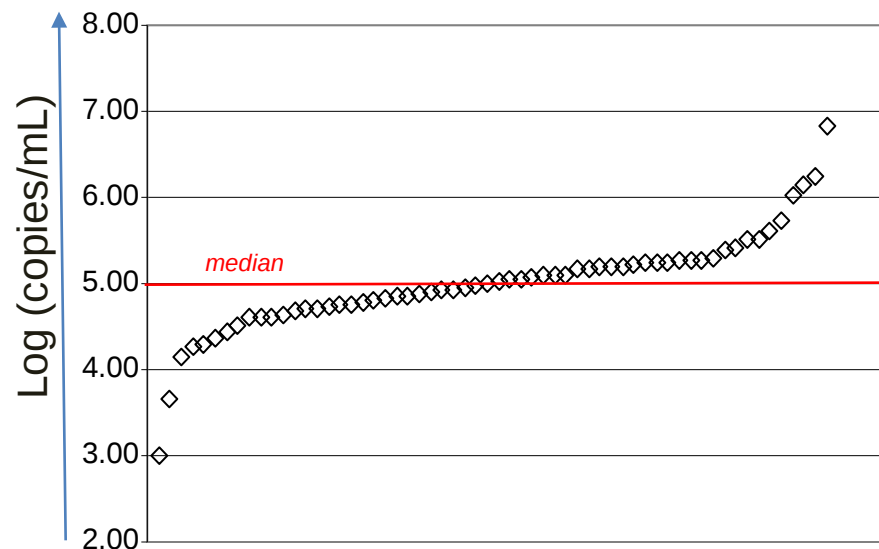
A total of 88 sets of specimens were distributed with 74 participants returning results, 13 participants did not examine the specimens

→61 sets of results

Pilot / Results by specimen

SPECIMEN A

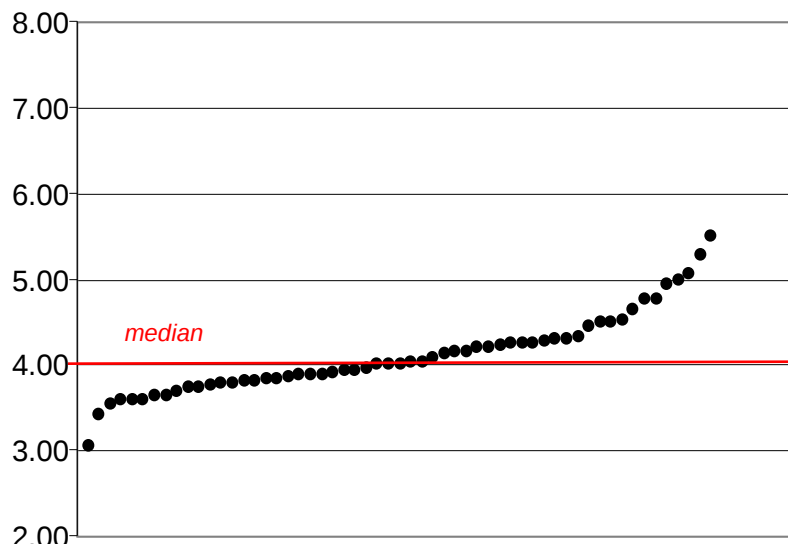
- Expected copies/mL 5.19 log (156,000)
- median+/-SD 5.01 +/- 0.57
- range 3.00-6.84



61 results sorted in increasing order

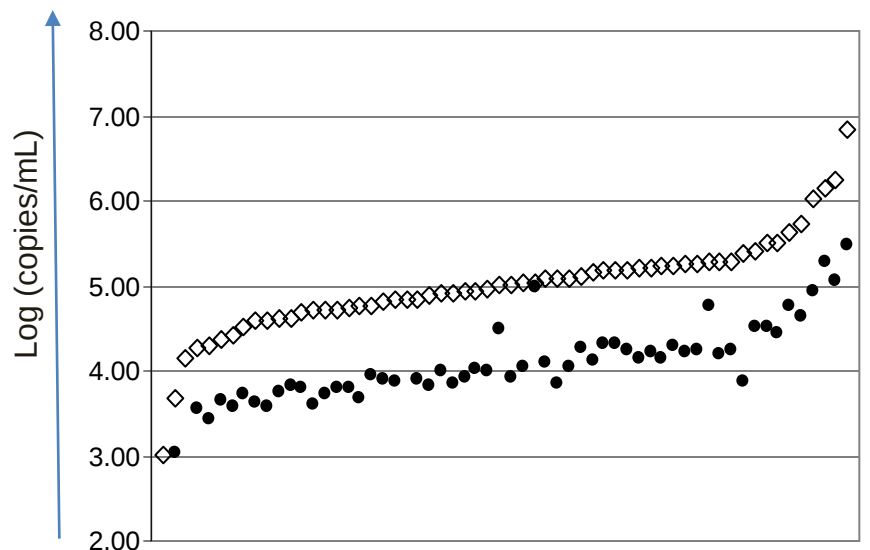
SPECIMEN B

- Expected copies/mL 4.19 log (15,600)
- median+/-SD 4.02 +/- 0.47
- range 3.05-5.49

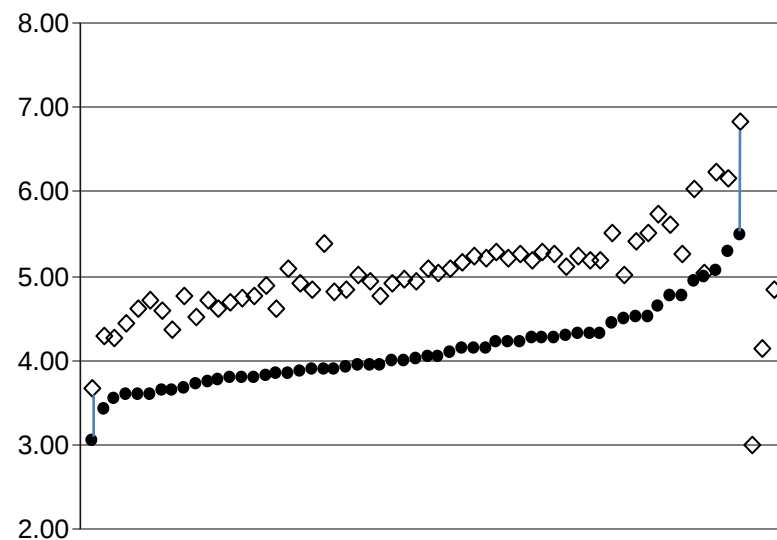


Pilot / Results by laboratory

- ◇ Specimen A
- Specimen B



Specimen-A results sorted
in increasing order



Specimen-B results sorted
in increasing order

Pilot / Results analysed as a log difference

98.4% (60/61) detected 5 log of EBV DNA (specimen A)

93.4% (57/61) detected 4 log of EBV DNA (specimen B)

-> 57 log differences calculated:

expected	1.00 log
median+/-SD	0.96+/-0.21
range	0.05 - 1.49

→ 96.5% (55/57) reported a 0.46 to 1.46 log difference

Pilot / Results analysed as a log difference

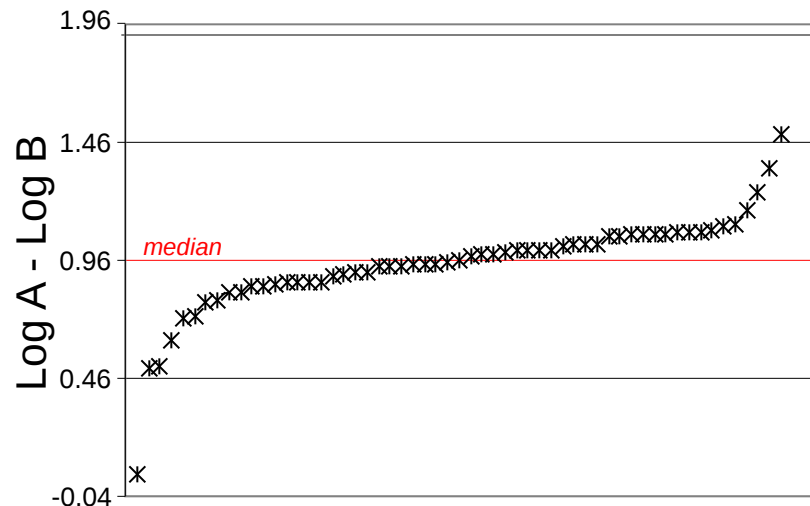
98.4% (60/61) detected 5 log of EBV DNA (specimen A)

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-> 57 log differences calculated:

expected	1.00 log
median+/-SD	0.96+/-0.21
range	0.05 - 1.49

→ 96.5% (55/57) reported a 0.46 to 1.46 log difference





- 1) same amplification assay
- 2) different amplification platforms
- 3) different extraction assays

Pilot / Extraction and amplification methods

Amplification method	n	%
Nanogen AD: Real-Time Alert	21	34.4%
Qiagen: Artus	17	27.9%
Real-Time Single target	14	23.0%
Roche: LightCycler	2	3.3%
Argene Biosoft	2	3.3%
PCR: Single target	1	1.6%
Real Time Multiplex	1	1.6%
Sacace Real-TM	1	1.6%
Unspecified	1	1.6%
Other	1	1.6%

- 26.2% in house assays
- 73.8% commercial assays
- All open systems:
free choice of extraction methods:

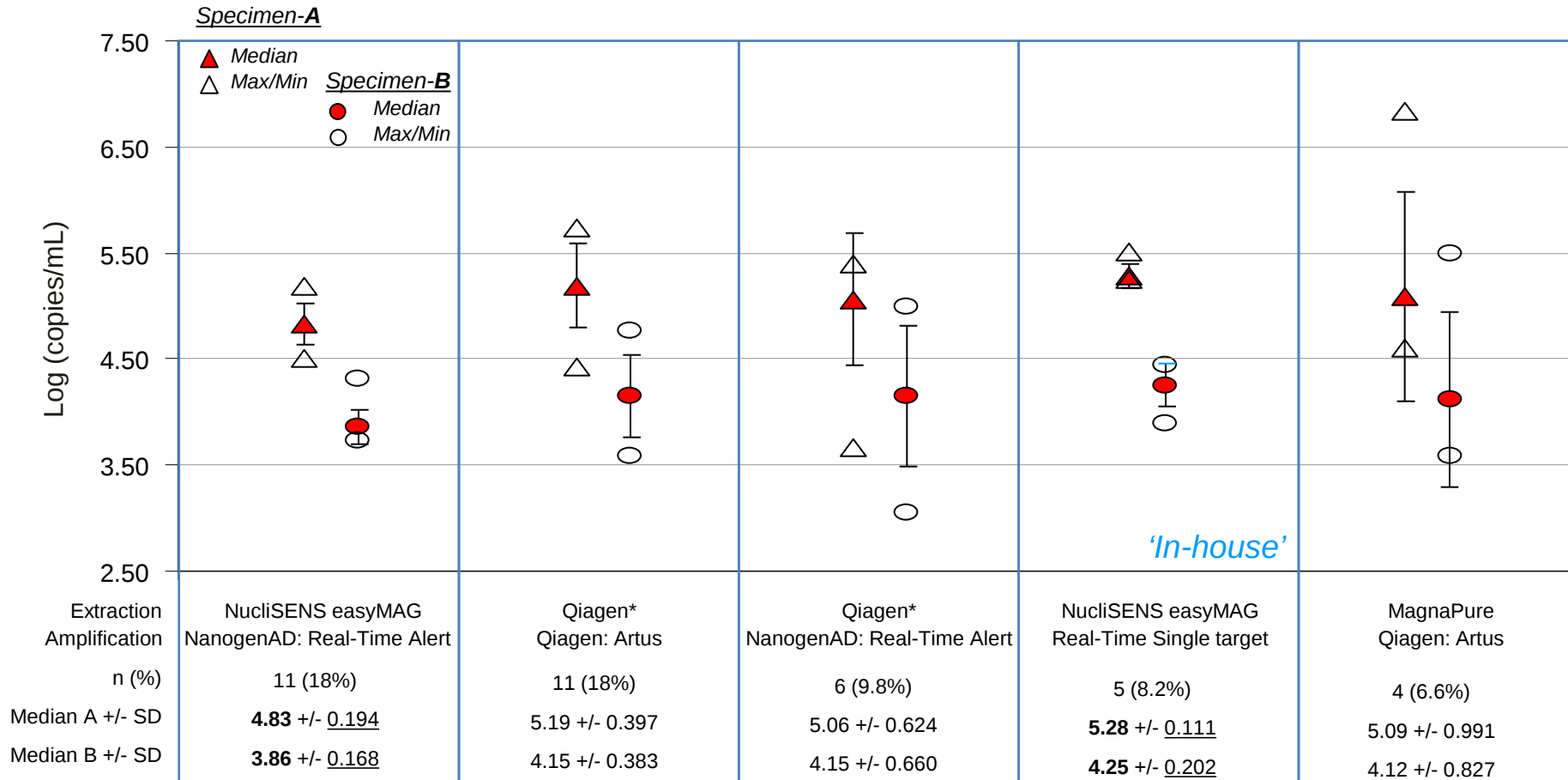
Extraction method	n	%
Qiagen*	24	36.9%
NucliSENS easyMAG	19	29.2%
MagnaPure	9	13.8%
Magnetic beads	2	3.1%
Nucleo Spin	1	1.5%
Unspecified	3	4.6%
Other	3	4.6%

* BioRobot MDx QIAasympohony

- >23 combinations of extraction and amplification assays
- 5 most frequent combinations:

Extraction method	Amplification method	n	%
NucliSENS easyMAG	Nanogen AD: Real-Time Alert	11	18.0
Qiagen*	Qiagen: Artus	11	18.0
Qiagen*	Nanogen AD: Real-Time Alert	6	9.8
NucliSENS easyMAG	Real-Time Single target	5	8.2
MagnaPure	Qiagen: Artus	4	6.6

Pilot / Results for the 5 most frequent E&A assay combinations



*BioRobot MDx, QIAasymphony

Conclusion

- This pilot study illustrates the lack of standardisation, with participants results ranging 3.84 log for specimen A (5 log) and 2.44 log for specimen B (4 log) using different commercial and in house assays.
- Standardisation:
 - o Matrix
 - o Extraction assay
 - o Amplification platform
 - o Amplification/detection assay
- Needed to establish guidelines for the surveillance of HSCT and SOT at risk of developing PTLD:
 - o Clinical cut-off for pre-emptive therapy
 - o Frequency of assessment (+)
 - o Treatment efficacy

EBV DNA quantification scheme

- To assess testing laboratories performance in the measurement of the EBV viral load
- Sample format: Freeze-dried plasma
- No. of distributions / year = 3
- No. of samples / distribution = 2
- Scoring: reported log difference in viral load between specimen pair
- Absolute quantitation data by method is also presented in the scheme distribution reports

EBV DNA quantification scheme

- Virus source: Namalwa cells
- Plasma EBV VCA IgG and EBNA G antibody status tested
- Number data sets 60
- No. method groups 10
- No. commercial 7
- No. commercial with >5 participants
 - Qiagen Artus, Artus Biosoft, ELITech: ELITe MGB and Nanogen AD RT Alert
- 3 participants stated their method was standardised against the IS
 - 1 gave a conversion factor of 13 (in house RT single target)

EBV: Current number of users by method

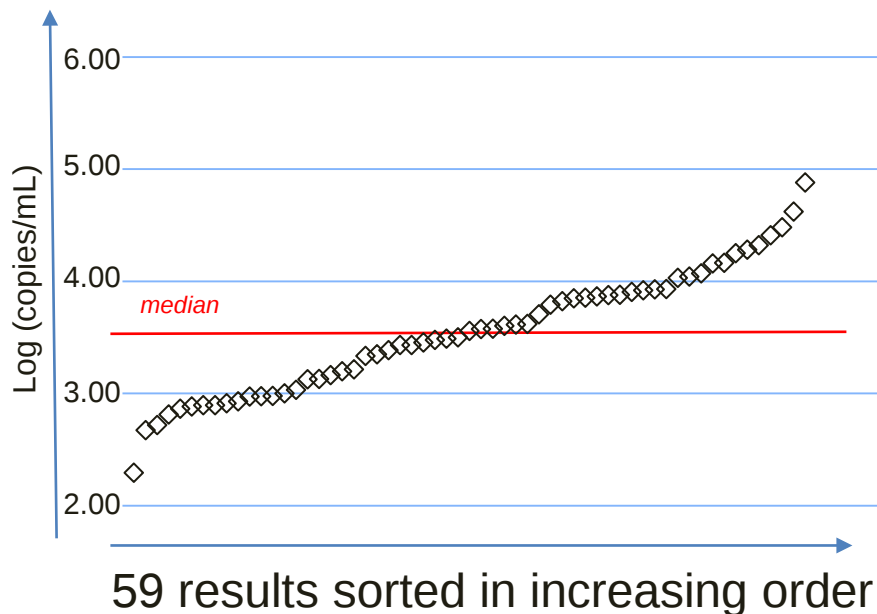
Unspecified	1
Focus: Simplexa	1
FTD	1
Altona: RealStar	2
Nanogen AD: R-T Alert	5
Argene Biosoft	6
ELITech: ELITe MGB	7
Real-Time Multiplex	8
Real-Time Single target	11
Qiagen: Artus	16

Distribution number: 3683

Results by specimen

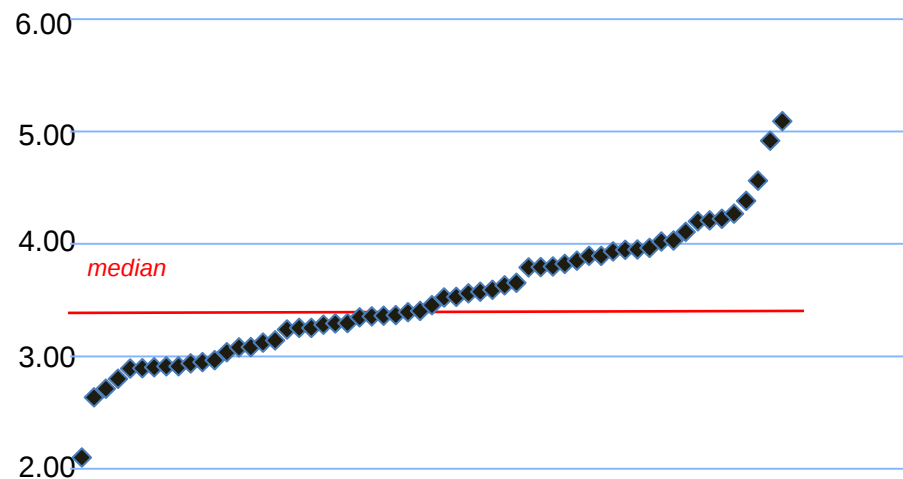
SPECIMEN 2620

- Expected copies/mL 3.47 log (2,900)
- median+/-SD 3.56 +/- 0.54
- range 2.29-4.88



SPECIMEN 2621

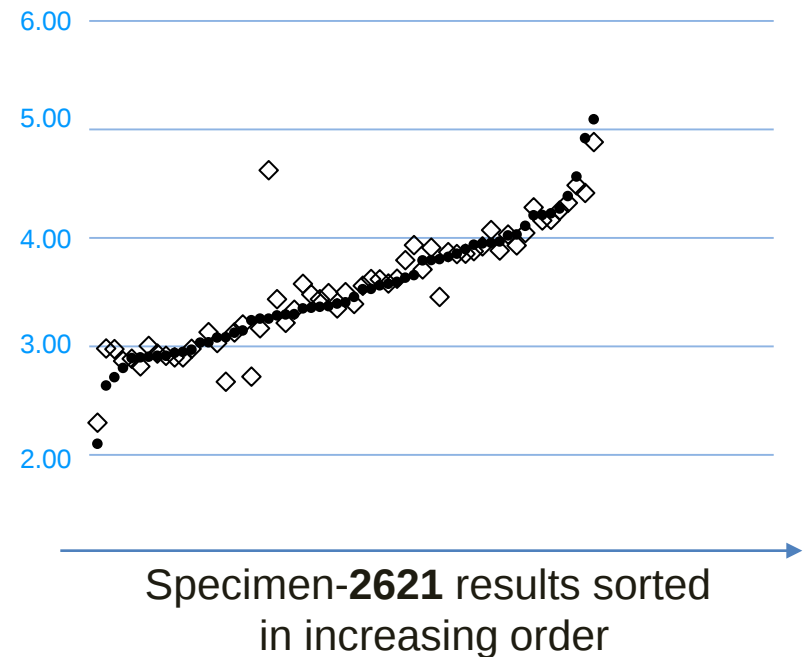
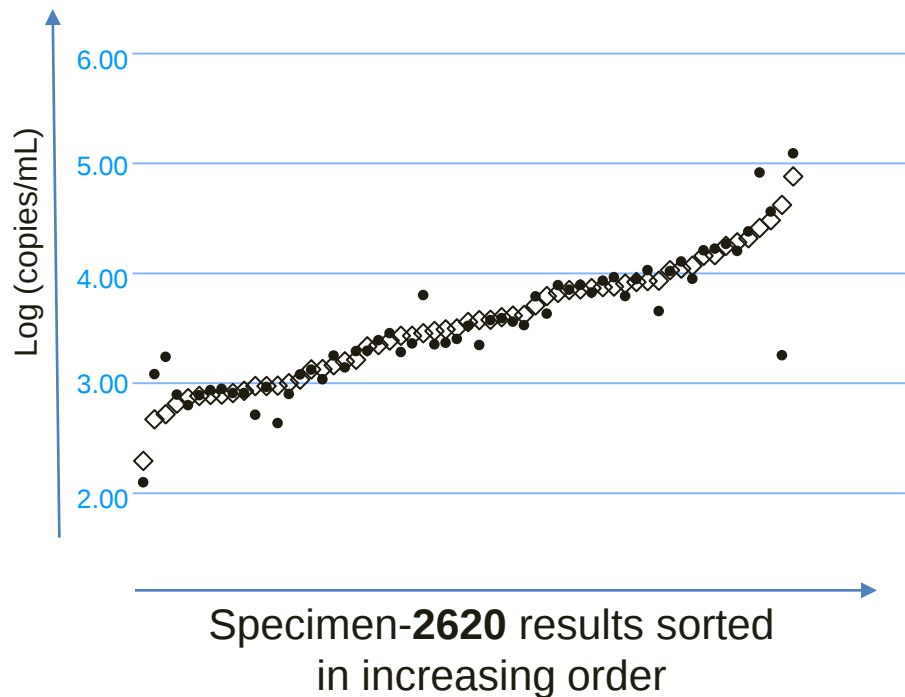
- Expected copies/mL 3.47 log (2,900)
- median+/-SD 3.43 +/- 0.57
- range 2.09-5.09



Distribution number: 3683

Results by laboratory

- ◇ Specimen 2620
- Specimen 2621



Results analysed as a log difference

96.7% (59/61) detected 3.5 log of EBV DNA (specimen 2620)

96.7% (59/61) detected 3.5 log of EBV DNA (specimen 2621)

-> 59 log differences calculated:

expected	0.00 log
median+/-SD	0.00+/-0.28
range	-0.52 - 1.37

→ 96.6% (57/59) reported a -0.51 to 0.49 log difference

Results analysed as a log difference

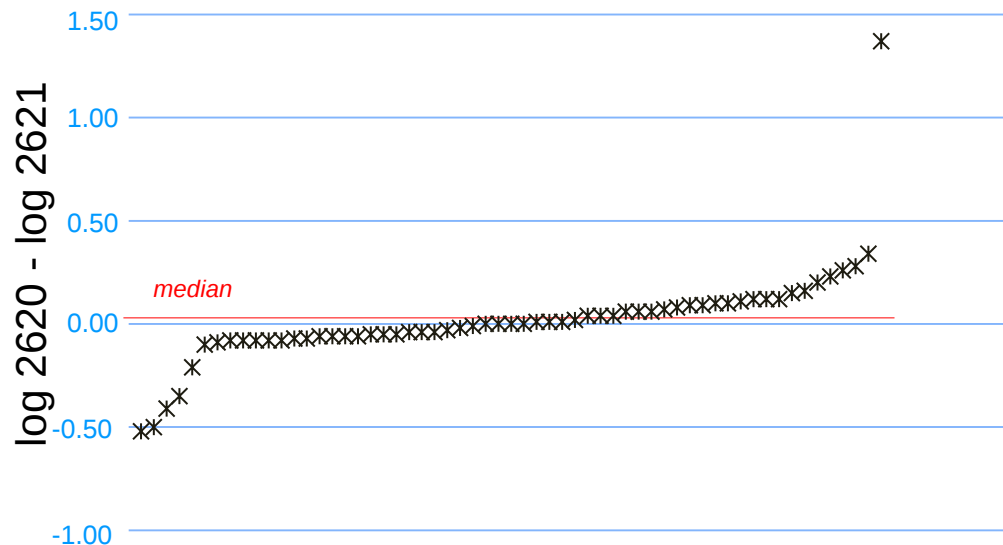
96.7% (59/61) detected 3.5 log of EBV DNA (specimen 2620)

96.7% (59/61) detected 3.5 log of EBV DNA (specimen 2621)

-> 59 log differences calculated:

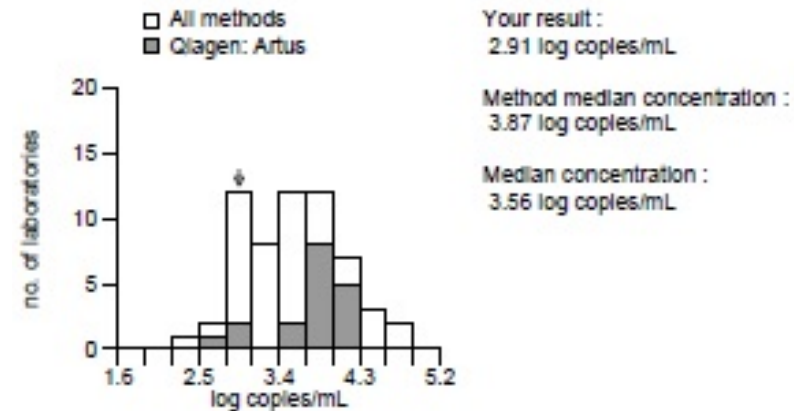
expected	0.00 log
median+/-SD	0.00+/-0.28
range	-0.52 - 1.37

→ 96.6% (57/59) reported a -0.51 to 0.49 log difference

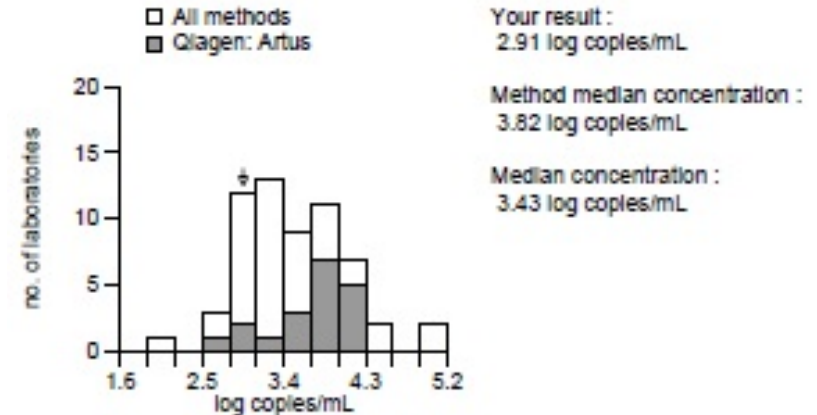


Distribution number: 3683

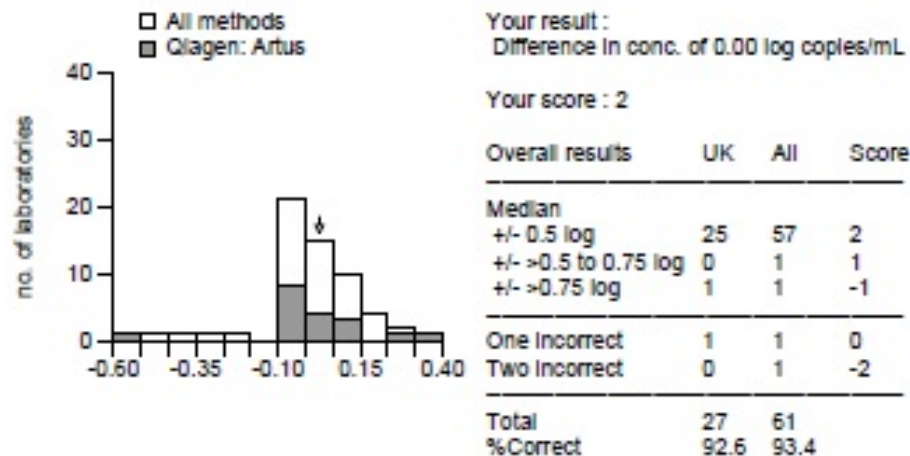
Specimen: 2620



Specimen: 2621

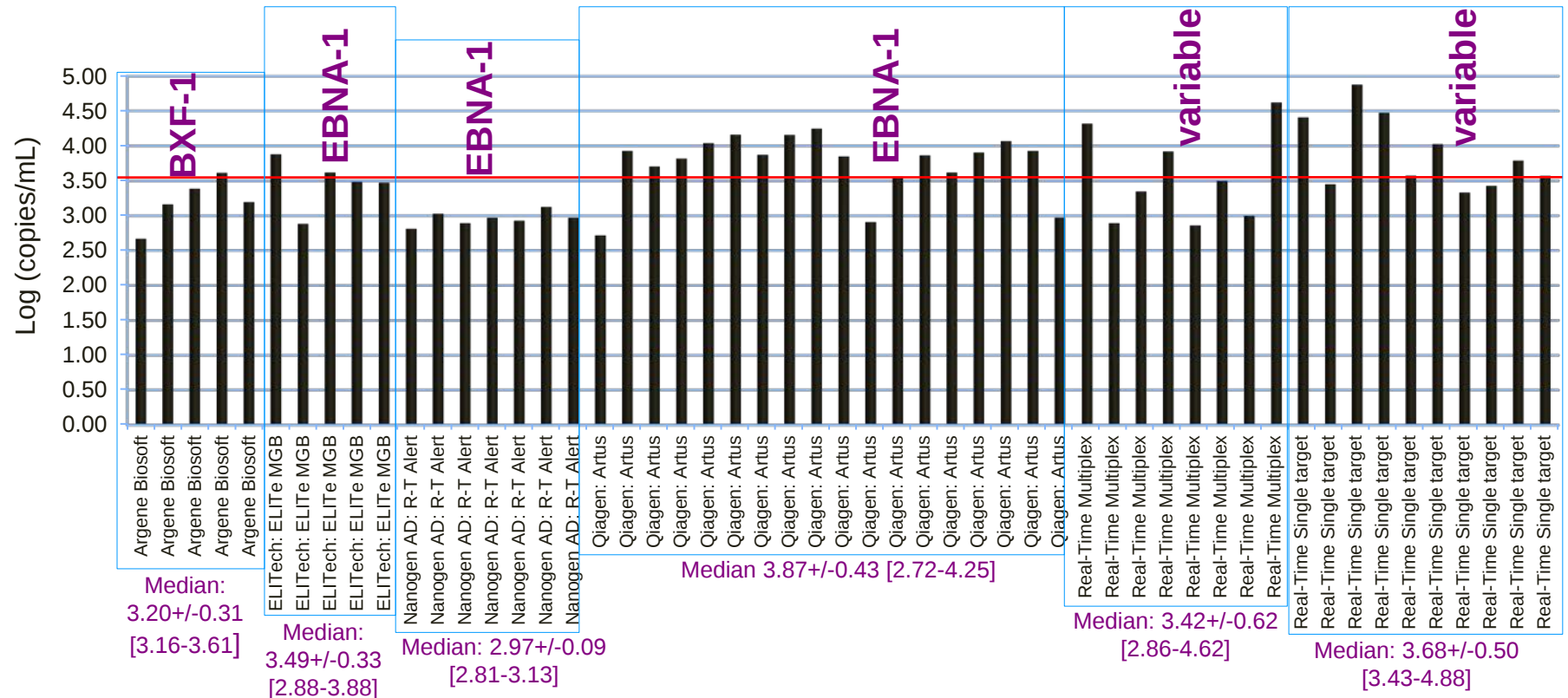


Intended result: -0.50-0.50 log copies/mL



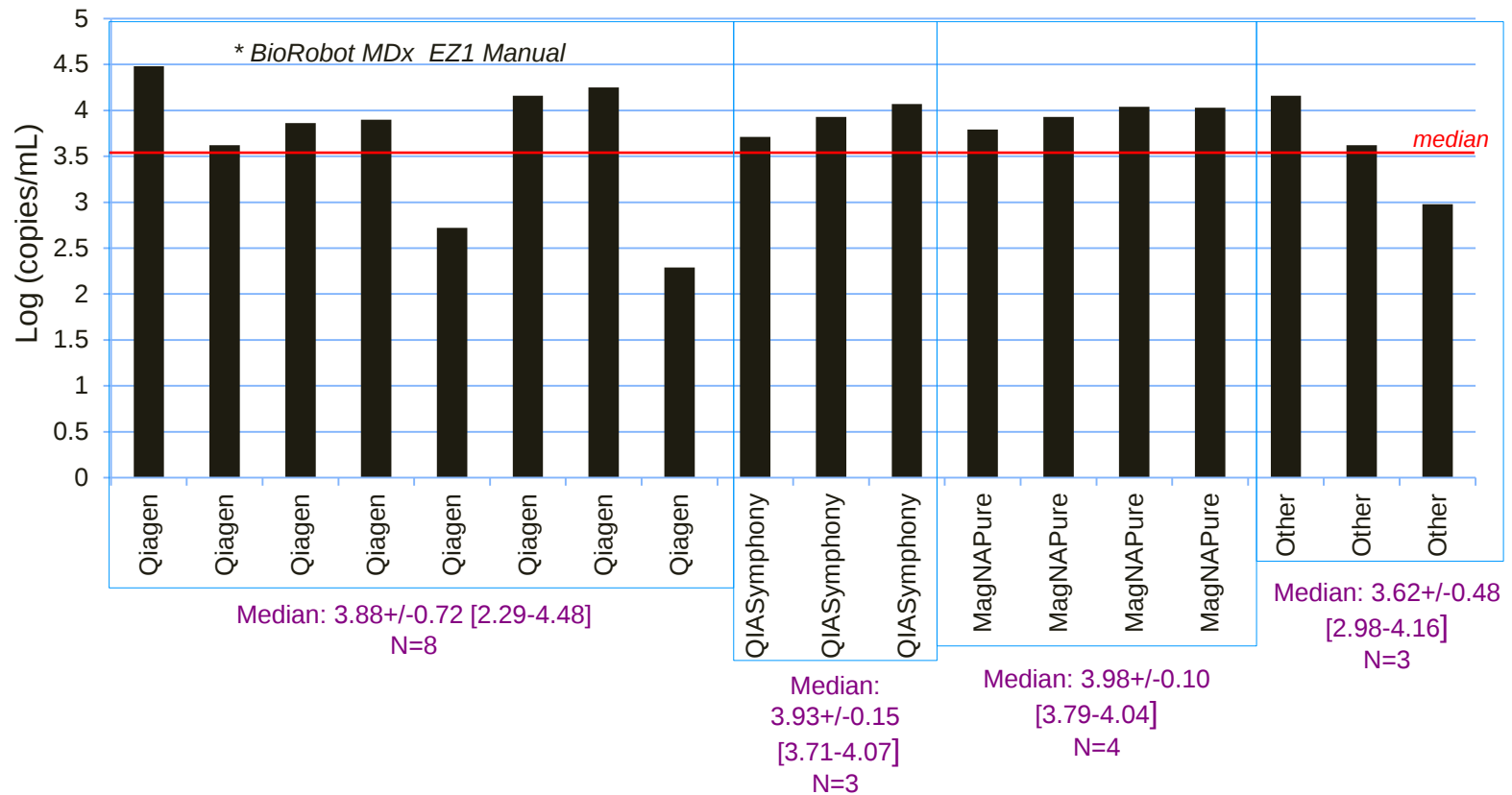
Specimen number: 2620

Results by amplification methods



Specimen number: 2620

Qiagen: Artus results by extraction methods



Extraction and amplification methods

Amplification method	N pilot	% pilot	N now	% now
Nanogen AD: Real-Time Alert	21	34.4	8	11.8
QiaGen: Artus	17	27.9	20	29.4
Real-Time Single target	14	23.0	11	16.2
Roche: LightCycler	2	3.3	0	0
Argene Biosoft	2	3.3	5	7.4
PCR: Single target	1	1.6	0	0
Real Time Multiplex	1	1.6	9	13.2
Sacace Real-TM	1	1.6	0	0
ELITech: ELITE MGB	0	0	9	13.2
Unspecified	1	1.6	1	1.5
Other	1	1.6	5	7.4

* BioRobot MDx QIA Symphony EZ1

• 29.4% in house assays (in pilot 26.2%)

• 70.6% commercial assays: all use IC

• Free choice of extraction methods:

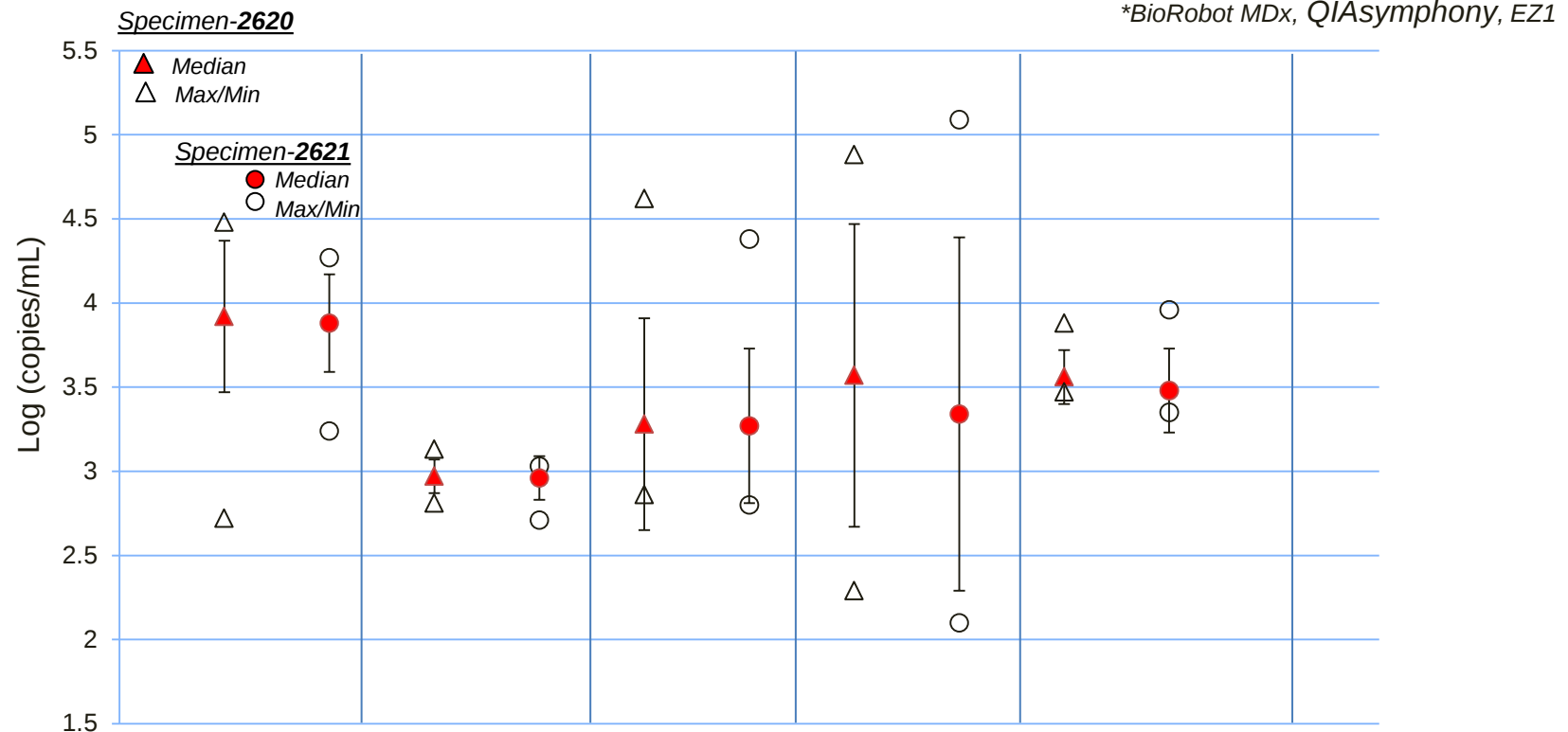
Extraction method	N Pilot	% Pilot	N now	% now
QiaGen*	24	36.9	31	51.7
NucliSENS easyMAG	19	29.2	18	30
MagnaPure	9	13.8	9	15
Magnetic beads	2	3.1	0	0
Nucleo Spin	1	1.5	1	1.7
Abbott: m2000sp	0	0	1	1.7
Unspecified	3	4.6	0	0
Other	3	4.6	0	0

- >22 combinations of extraction and amplification assays
- 5 most frequent combinations

Extraction method	Amplification method	N Pilot	% Pilot	N Now	% Now
NucliSENS easyMAG	Nanogen AD: Real-Time Alert	11	18.0	6	11.3
QiaGen*	QiaGen: Artus	11	18.0	12	22.6
QiaGen*	Nanogen AD: Real-Time Alert	6	9.8	0	0
NucliSENS easyMAG	Real-Time Single target	5	8.2	3	5.7
MagnaPure	QiaGen: Artus	4	6.6	4	7.5
QiaGen*	Real-Time Multiplex	0	0	8	15
QiaGen*	Real-Time Single target	0	0	5	9.4
NucliSENS easyMAG	ELITech: ELITE MGB	0	0	5	9.4

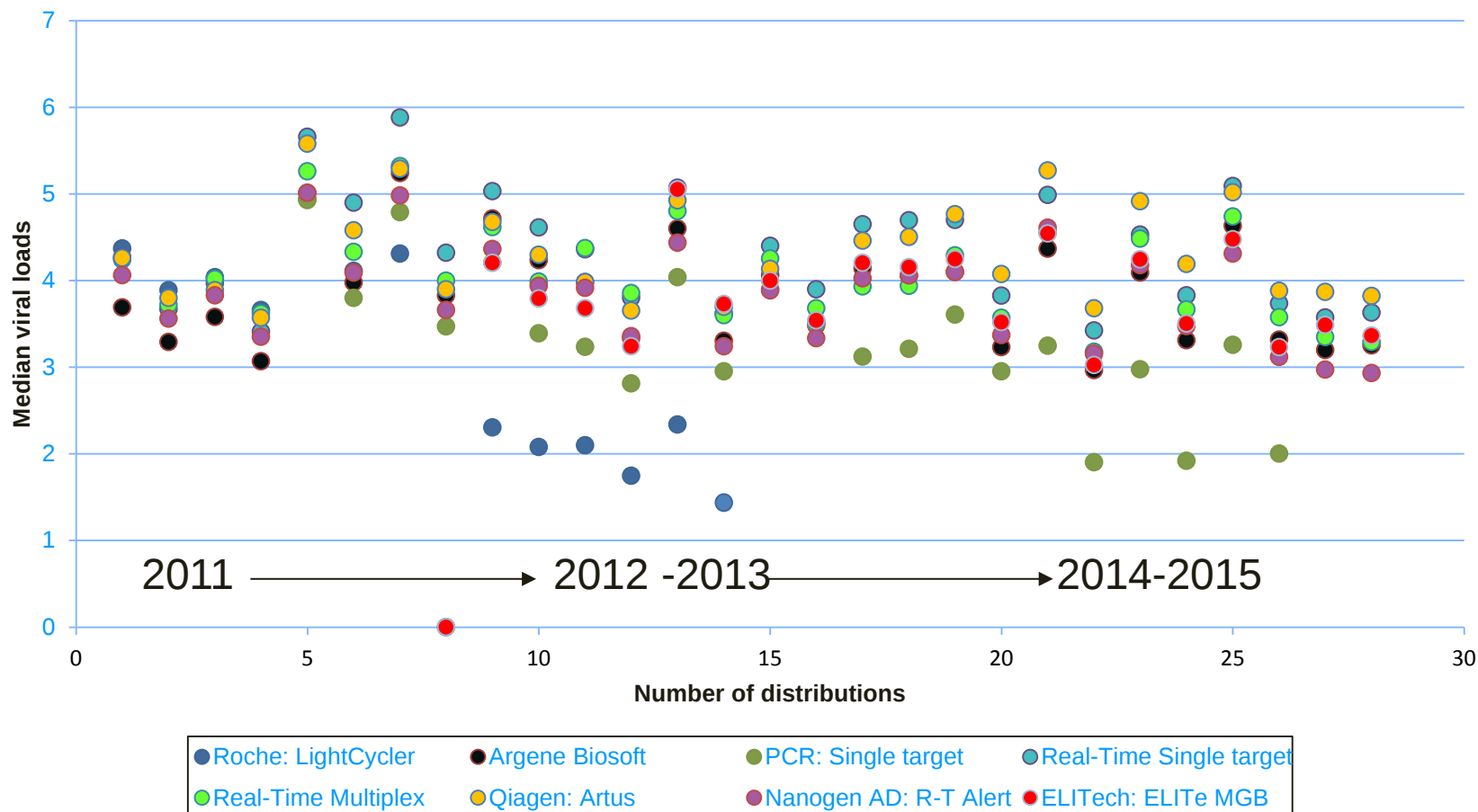
Distribution number 3683

Results for the 5 most frequent E&A assay combinations



Extraction	Qiagen*	NucliSENS easyMAG	Qiagen*	Qiagen*	NucliSENS easyMAG
Amplification	Qiagen: Artus	NanogenAD: Real-Time Alert	Real-Time Multiplex	Real-Time Single target	ELITech: ELITe MGB
n (%)	12 (20%)	6 (10%)	8 (13.3%)	5 (8.3%)	5 (8.3%)
Median 2620 +/- SD	3.92 +/- 0.45	2.97 +/- 0.10	3.28 +/- 0.63	3.57 +/- 0.90	3.56 +/- 0.16
Median 2621 +/- SD	3.88 +/- 0.29	2.96 +/- 0.13	3.27 +/- 0.46	3.34 +/- 1.05	3.48 +/- 0.25

EBV: Median viral load by method



Summary

- The introduction of the EBV WHO International Standard is relatively recent: use has not been widely adopted
- Several participants indicated they are in the process of introducing the Standard
- It is too soon to see any improvement in the comparability of results with different kits
- Reportable ranges
- Sample type
- Converting viral load values from c/mL to IU

Thank you for listening

Many thanks to:

- Colleagues from the Microbiology Laboratory at the Northern General Hospital, Sheffield for their kind assistance with pre-distribution testing
- Participants for their data
- Great team at UK NEQAS

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Vivienne James
and Maila Mutso