



Protecting and improving the nation's health

Summary of Results

Norovirus and Hepatitis A virus Scheme

External Quality Assessment for Microbiology

Distribution Number: NHV001

Sample Numbers: NH' J007 & NH v 0008

| Distribution Date: | 1 Ocւ ber 2018 |
|--|---------------------------------|
| Results Due: | : November 2018 |
| Report Date: | 12 November 2018 |
| Samples preruped and quality control tested by | Justin Avant (Cefas) |
| Data a Plys d by: | Nita Patel Manchari Rajkumar |
| l eport compiled by: | Nita Patel Manchari Rajkumar |
| Authorised by: | Nita Patel |

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Lab no:

The data in FEPTU reports is confidential

Overview:

This Scheme provides external quality assessment samples for laboratories that examine food products or waters for hepatitis A virus and norovirus using the reverse-transcription polymerase chain reaction (RT-PCR). Regulation (EC) No 669/2009 sets out specifications for an increased level of official controls on imports of certain feed and food of non-animal origin and via Implementing Regulation (EU) 2016/2107 currently applies to frozen raspberries imported from Serbia.

According to the European Food Safety Authority (EFSA) foodborne viruses are the second most common cause of foodborne outbreaks in the European Union (EU) after Salmonella https://www.food.gov.uk/science/microbiology/norovirus/norovirus-eu. EFSA has identified that removal of viral contamination is extremely difficult. It therefore recommends that the focus on control of viruses in the food chain needs to be based on preventing contamination and cross-contamination of food. Viruses cannot be cultured therefore molecular techniques are the methods of choice for the detection, identification and quantification of foodborne viruses.

ISO 15216-1:2017 Microbiology of the food chain -- Horizontal method for determination of hepatitis A virus and norovirus using real-time RT-PCR -- Part 1: Method for quantification is used by many laboratories. This ISO method covers pre-treatment steps to elute viruses from the different matrices.

This proficiency testing scheme challenges laboratories in detection and quantification (copies or comple) of hepatitis A virus (HAV) and Norovirus GI and GII. It has been organised in collaboration with Cefas as the European Union Reference Laboratory (EURL) for monitoring bacteriological and viral contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and viral contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and viral contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and viral contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and viral contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and Viral Contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and Viral Contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and Viral Contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and Viral Contamination of bivelenges as the European Union Reference Laboratory (EURL) for monitoring bacteriological and Viral Contamination (EURL) for monitoring bacteriological (EURL) for monitoring bacteriological (EURL) for monitoring bact

FEPTU Quality Control:

The samples were prepared and quality controlled at Cefas.

To demonstrate homogeneity of the sample 20 LENTICULE® discs selected and or in from a batch were examined.

To demonstrate stability of the sample a minimum of Ten LF' , FICU .E discs, selected randomly from a batch, were examined throughout the distribution period.

Cefas used a qRT-PCR using the RNA UltraSense™ One-Step Quantitative RT-PCR System (ThermoFisher), on a Stratagene Mx3005P real-time PCR machine.

The intended results letters provide guidance for participant, rear ding the intended result.

Guidelines and general advise:

If you experience difficulties with any of the ex mir. Ons please refer to section 17.0 of the Scheme Guide https://www.gov.uk/government/publicatio bod-ar war oroficiency-testing-scheme-scheme-guide

Please contact FEPTU staff for advive and information:

N.ta Patel

Repeat samples Came Comes or Kermin Daruwalla Tel: +44 (0)20 8327 7119

Data analysis lita From Manchari Rajkumar Fax: +44 (0)20 8200 8264

Microbiological 2 .vice | na . atel E-mail: foodeqa@phe.gov.uk

General comments and

complaints
Scheme Advisors

Justin Avantⁱ & James Lowtherⁱ

Scheme Co-ordinator Nita Patel

Accreditation:

PHE will be applying for this scheme to be accredited with the United Kingdom Accreditation Service (UKAS) to ISO/IEC 17043:2010

FEPTU's website

ⁱThe Centre for Environment, Fisheries and Aquaculture Science, European Union Reference Laboratory for monitoring bacteriological and viral contamination of bivalve molluscs, Weymouth Laboratory, Dorset, DT4 8UB, United Kingdom

| Total number of participants sent distribution NHV004 | 36 |
|--|----|
| Number of laboratories not returning a result for NHV004 | 1 |
| Number of laboratories not examining any samples in NHV004 | 1 |

Sample: NHV0007

Sample type: LENTICULE $^{\circledR}$ discs prepared with known levels of norovirus GII from human faeces and known levels of Hepatitis A virus (HAV) cell culture supernatant.

Request:

Examine for the presence of viruses (norovirus GI and GII, hepatitis A virus) Quantify these viruses in the sample (if routinely done)

Contents and summary of results:

| Examination | on | Expected result | Your Result | PHE Score | Z-score |
|-------------|---------------|-----------------|-------------|-----------|---------|
| | Norovirus GI | Positive | | | |
| Virus | Norovirus GII | Positive | | | |
| | HAV | Negative | | | |

Quantification results

| Examinati | ion | Expected range | Your Result | PHF ≏or€ | Z-score |
|-----------|---------------|--------------------------|-------------|----------|---------|
| | Norovirus GI | $60 - 2.5 \times 10^3$ | | | |
| Virus | Norovirus GII | 58 – 3.6x10 ³ | | R | |
| | HAV | - | | | |

Norovirus GI (graph of results shown on page 4)

| Total participants reporting for Norovirus GI | 32 |
|---|--|
| Participants reporting correctly a detected result | 30 (94%) |
| Number of laboratories reporting a result copie , per , mp e | 15 |
| Expected range as copies per sample | $60 - 2.5 \times 10^3 (1.78 \log_{10} - 3.39 \log_{10})$ |
| Assigned value (participants' median' | 3.8x10 ² copies per sample (2.58 log ₁₀) |
| Uncertainty of assigned value ($U(X_{\rho_{\nu}}) = \log_{10}$ | 0.13 |
| No. of outlying counts | 1 (1 high) |
| Participants mean | 3.6x10 ³ copies per sample (3.55 log ₁₀) |
| Standard devi non of par sipents results ** | 0.40 copies per sample log ₁₀ |
| FEPTU's Comercian | 1.70x10 ² copies per sample (2.23 log ₁₀) |

Norovirus GII (پر یا h of results shown on page 4)

| Total participants reporting for Norovirus GII | 32 |
|---|--|
| Participants reporting correctly a detected result | 28 (88%) |
| Number of laboratories reporting a result copies per sample | 13 |
| Expected range as copies per sample | $58 - 3.6 \times 10^3 (1.76 \log_{10} - 3.56 \log_{10})$ |
| Assigned value (participants' median) | 4.6x10 ² copies per sample (2.66 log ₁₀) |
| Uncertainty of assigned value ($U(X_{pt}) = \log_{10}$) | 0.16 |
| Participants mean | 5.51x10 ² copies per sample (2.74 log ₁₀) |
| Standard deviation of participants results ** | 0.45 copies per sample log ₁₀ |
| FEPTU's QC median | 4.5x10 ² copies per sample (2.65 log ₁₀) |

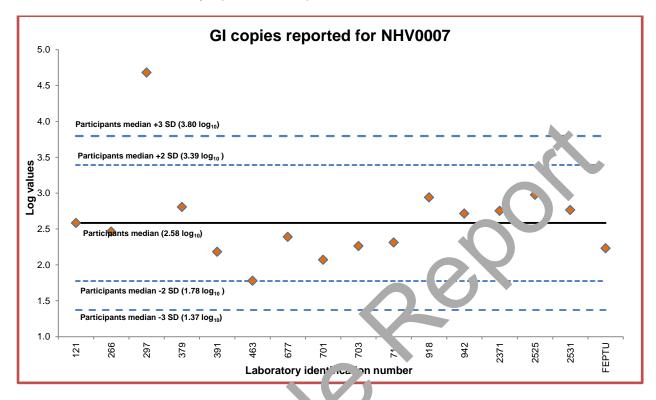
^{**} Robust SD based on median absolute deviation about the participants' median (MADe)

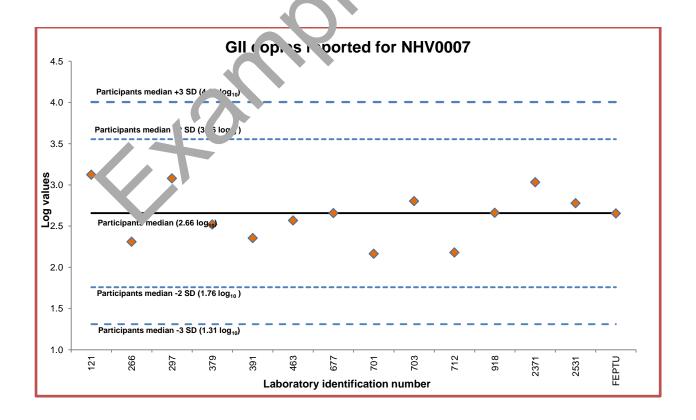
HAV

| Total participants reporting for HAV | 33 |
|--|----------|
| Participants reporting correctly reporting a not detected result | 32 (97%) |

Sample specific comment

Two laboratories reported a false negative result for Norovirus GI, four laboratories reported a false negative result for GII and one laboratory reported a false positive result for HAV.





Sample: NHV0008

Sample type: $LENTICULE^{@}$ discs prepared with known levels of norovirus GII from human faeces and known levels of Hepatitis A virus (HAV) cell culture supernatant.

Request:

Examine for the presence of viruses (norovirus GI and GII, hepatitis A virus) Quantify these viruses in the sample (if routinely done)

Contents and summary of results:

| Examir | nation | Expected result | Your Result | PHE Score | Z-score |
|--------|---------------|-----------------|-------------|-----------|---------|
| | Norovirus GI | Negative | | | |
| Virus | Norovirus GII | Negative | | | |
| | HAV | Positive | | | |

Quantification results

| Examinati | on | Expected range | Your Result P. 'E Scr | re Z-score |
|-----------|---------------|-----------------------|-----------------------|------------|
| | Norovirus GI | - | | |
| Virus | Norovirus GII | - | 00, | |
| | HAV | $1.3x10^2 - 4.6x10^4$ | | |

Norovirus GI

| Total participants reporting for Norovirus GI | <u>_</u> | 32 |
|---|----------|----------|
| Participants reporting correctly a not detected esult | | 31 (97%) |

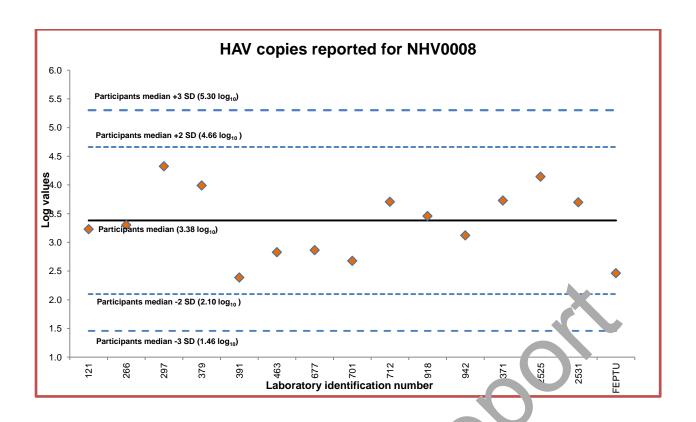
Norovirus GII

| Total participants reporting for Nor irus C' | 32 |
|--|----------|
| Participants reporting correctly a not de acted result | 31 (97%) |

HAV (graph of results show an page 6)

| Total participants reportination 1AV | 33 |
|--|--|
| Participants reprising correctly reporting detected result | 32 (97%) |
| Number of labo. tor s reporting a result copies per sample | 14 |
| Expected range as copies per sample | $1.3x10^2 - 4.6x10^4 (2.10 \log_{10} - 4.66 \log_{10})$ |
| Assigned value (participants' median) | 2.4x10 ³ copies per sample (3.38 log ₁₀) |
| Uncertainty of assigned value ($U(X_{pt}) = \log_{10}$) | 0.21 |
| Participants mean | 5.0x10 ³ copies per sample (3.70 log ₁₀) |
| Standard deviation of participants results ** | 0.64 copies per sample log ₁₀ |
| FEPTU's QC median | 2.90x10 ² copies per sample (2.46 log ₁₀) |

^{**} Robust SD based on median absolute deviation about the participants' median (MADe)



General comments for this distribution

This is the fourth distribution of the Norovirus and Hepatitis A virus scheme since it was launched in April 2017.

The results data suggests that one participant has either swapped their samples during testing or entered their results incorrectly.

Scoring applied

The samples in this distribution have been scored using the below scoring criteria.

Presence/absence results

Participants' correct results for detection are allocated scores up to a maximum of two points as follows:

| Fully correct result for the intended result | 2 |
|--|---|
| False positive / false negative result | 0 |

Quantification results

The expected range for each copies per sample result reported is calculated using the median absolute deviation from the median (MADe**) values (see * below) which are determined from the median result reported by participants' and take into account the following criteria:

- (1) median ± 2 MADeS*
- (2) median ± 3 MADeS*
- (3) median ± 0.5 log₁₀ units

If the ranges in (1) and/or (2) are less than the value of the r rediar $\pm 0.5 \log_{10} units$ then the expected range is extended as described in (3).

| | Score |
|--|-------|
| Expected range within the range according to criterir (1) | 2 |
| Outlying results (1) within the range of criteria (2, bu ot w hin criteria (1) | 1 |
| Outlying results (2) outside the range of criter (2) | 0 |

Non-return of results

Participants who do not return a result by the specified date are allocated a PHE score of zero for all tests.

*Statistics applied for copi ner ample

The median absolute deviation from the median (*MADe*) values is used to identify outlying counts when there are less than 50 of talse. The use of MAD values provided statistically robust method for calculating the acceptable range using an analysis the requires calculation of the median difference from the median for every participant's result. This is then multiplied by a constant (1.4826) to obtain a robust estimate of the standard deviation (MAD value). The results are initially interpreted as follows:

Results in range of participants' median ±2 MADSD satisfactory
Results between ±2 MAD and ±3MADSD questionable
Results >±3 MADSD unsatisfactory

Statistical evaluation for quantification results

Participants are advised that for a robust statistical evaluation for quantification data at least 20 results are required for each examination. When statistical calculation is based on 10 – 19 results, they should be interpreted with caution as they may be overly influenced by outlying results. When there are fewer than 10 reported results, the statistics are not considered robust to enable the examination to be scored.

| | Summary of participants results NHV0007 (any incorrect results are shown in red) | | | | | | | | | | | |
|-----|--|---------|-----------|----------|-----------|--------|---|--|--|--|--|--|
| Lab | Norov | irus GI | Norovi | irus GII | HA | AV | RNA extraction | | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | | |
| | + | 384 | + | 1326 | - | 0 | BioMérieux NucliSENS® miniMag® Inv rogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | | |
| | + | | + | | - | | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | | |
| | + | | + | | | | Qiagen QIAamp® Viral RNA CeeramTools® GI, GII and HAV detection kits | | | | | |
| | + | 291 | + | 204 | | | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | | |
| | + | | + | | | | In house CeeramTools® GI, GII and HAV detection kits | | | | | |
| | + | 48000 | + | 1200 | | | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | | |
| | + | | + | | - | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus/Hepatitis A 3plex kit | | | | | |
| | + | | + 0 | | - | | KingFisher™ MagMAX™ CORE Nucleic Acid Purification Kit Applied Biosystems® TaqMan® Universal PCR Master Mix | | | | | |
| | + | 640 | 7 | 330 | - | <20 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | | |
| | + | 152 | + | 226 | - | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | | |
| | + | | + | | - | | BioMérieux NucliSENS® miniMag® | | | | | |

| | Summary of participants results NHV0007 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|---------|-----------|---|-----------|--------------|---|--|--|--|--|
| Lab | Norov | irus GI | Norovi | rus GII | H | AV | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | | | | | | | QIAGEN® QuantiTect® Probe RT-PCR kit | | | | |
| | + | | + | | - | | BioMérieux NucliSENS® miniMag® In itroyen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | 60 | + | 369 | | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | | + | | | | Stratec INVISORB Spin Virus RNA Mini Kit Altona DIAGNOSTICS RealStar® Norovirus RT-PCR Kit 2.0 | | | | |
| | + | 245 | + | 454 | 0 | Not detected | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | 118 | + | 146 | | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | 183 | + | 636 | - | Not detected | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | 205 | 10 | 151 | - | | QIAGEN QIAsymphony DSP virus/pathogen mini kit Invitrogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | |
| | + | 872 | + | 457 | - | - | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | + | 519 | + | <loq< td=""><td>-</td><td></td><td>Roche High Pure Viral RNA kit Applied Biosystems™ TaqMan™ Fast Virus One-Step Master Mix</td></loq<> | - | | Roche High Pure Viral RNA kit Applied Biosystems™ TaqMan™ Fast Virus One-Step Master Mix | | | | |

| | Summary of participants results NHV0007 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|---------|--------------|---------------|-------------------|--------|--|--|--|--|--|
| Lab | Norovi | irus GI | Norovi | virus GII HAV | | AV | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | + | | + | | | | QIAGEN QIAamp® Viral RNA CeeramTools® GI, GII detection kits | | | | |
| | + | | + | | - | | CONGEN SureFast® Prep DNA/RNA virus | | | | |
| | Not examined | | Not examined | | | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | | + | | | | QIAGEN RNeasy kit CeeramTools® GI, GII and HAV detection kits | | | | |
| | | | | | Non-return of res | sults | | | | | |
| | + | | + | | | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | | + | | - | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | + | | :0 | | - | | BioMérieux NucliSENS® miniMag® Applied Biosystems™ AgPath-ID™ One-Step PCR Reagents | | | | |
| | + | 568 | 7 | 1075 | - | | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | 0 | | 0 | + | 3210 | MACHEREY-NAGEL NucleoSpin® RNA Virus ISO 15216-1:2017MasterMix Invitrogen | | | | |
| | + | | + | | - | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus/Hepatitis A 3plex kit | | | | |

| | Summary of participants results NHV0007 (any incorrect results are shown in red) | | | | | | | | | | | |
|-----|--|---------|-----------|---------|---------------|--------|---|--|----------------|--|--|--|
| Lab | Norov | irus GI | Norovi | rus GII | HAV | | HAV | | RNA extraction | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | | |
| | | | | | - | | BioMérieux NucliSENS® miniMag® Inv rogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | | |
| | + | 947 | - | | - | | QIAGEN QIAamp® Viral RNA QIAGEN OneStep RT-PCR kit | | | | | |
| | + | 580 | + | 600 | | 0 | BIOTECON Diagnostics Extraction Kit | | | | | |
| | - | | - | | Not exa. inc. | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus PLUS | | | | | |
| | Not examined | | | | 0 | | BIOTECON Diagnostics Extraction Kit | | | | | |

| | Summary of participants results NHV0008 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|---------|-----------|---------|------------|--------|---|--|--|--|--|
| Lab | Norov | irus GI | Norovi | rus GII | HAV | | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | - | 0 | - | 0 | + | 1702 | BioMérieux NucliSENS® miniMag® Inv rogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | |
| | - | | - | | + | | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | | - | | + | (2) | Qiagen QIAamp® Viral RNA CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | | - | | + | 2008 | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | | - | | C † | NE | In house CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | | - | | + | 21200 | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | - | - | | + | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus/Hepatitis A 3plex kit | | | | |
| | - | | . 0 | | + | | KingFisher™ MagMAX™ CORE Nucleic Acid Purification Kit Applied Biosystems® TaqMan® Universal PCR Master Mix | | | | |
| | - | <20 | , + | <20 | + | 9800 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | | | + | 244 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | | + | | BioMérieux NucliSENS® miniMag® | | | | |

| | Summary of participants results NHV0008 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|--------------|-----------|--------------|-----------|--------|---|--|--|--|--|
| Lab | Norov | irus GI | Norovi | irus GII | H | AV | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | | | | | | | QIAGEN® QuantiTect® Probe RT-PCR kit | | | | |
| | - | | - | | + | | BioMérieux NucliSENS® miniMag® In itroyen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | | + | 674 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | | 0 | | Stratec INVISORB Spin Virus RNA Mini Kit Altona DIAGNOSTICS RealStar® Norovirus RT-PCR Kit 2.0 | | | | |
| | - | Not detected | - | Not detected | | 733 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | | + | 478 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | Not detected | - | No 1etected | + | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | 1,5 | | + | 5080 | QIAGEN QIAsymphony DSP virus/pathogen mini kit Invitrogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | |
| | - | - | | - | + | 2880 | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | - | | - | | + | 1322 | Roche High Pure Viral RNA kit Applied Biosystems™ TaqMan™ Fast Virus One-Step Master Mix | | | | |

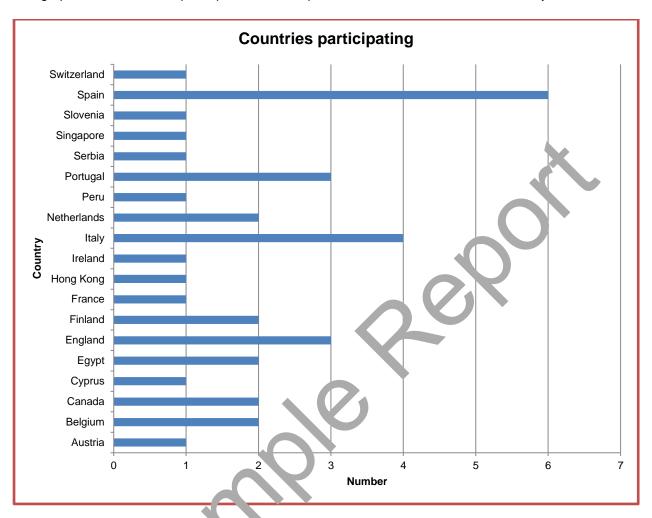
| | Summary of participants results NHV0008 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|---------|--------------|----------|-------------------|--------|--|--|--|--|--|
| Lab | Norov | irus GI | Norovi | irus GII | GII HAV | | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | - | | - | | | | QIAGEN QIAamp® Viral RNA CeeramTools® GI, GII detection kits | | | | |
| | - | | - | | + | | CONGEN SureFast® Prep DNA/RNA virus | | | | |
| | Not examined | | Not examined | | + | 07 | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | | + | | QIAGEN RNeasy kit CeeramTools® GI, GII and HAV detection kits | | | | |
| | | | | | Non-return of res | sults | | | | | |
| | - | | - | | 0 | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | - | A P | + | | BioMérieux NucliSENS® miniMag® Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | | | | |
| | - | | :0 | | + | | BioMérieux NucliSENS® miniMag® Applied Biosystems™ AgPath-ID™ One-Step PCR Reagents | | | | |
| | - | | 7 | | + | 5361 | BioMérieux NucliSENS® miniMag® CeeramTools® GI, GII and HAV detection kits | | | | |
| | + | 946 | + | 891 | - | 0 | MACHEREY-NAGEL NucleoSpin® RNA Virus ISO 15216-1:2017MasterMix Invitrogen | | | | |
| | - | | - | | + | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus/Hepatitis A 3plex kit | | | | |

| | Summary of participants results NHV0008 (any incorrect results are shown in red) | | | | | | | | | | |
|-----|--|--------|-----------|---------|----------------|-------------|--|--|--|--|--|
| Lab | Norovi | rus GI | Norovi | rus GII | HAV | | RNA extraction | | | | |
| | Detection | Copies | Detection | Copies | Detection | Copies | RT PCR reagent | | | | |
| | | | | | + | | BioMérieux NucliSENS® miniMag® Inv. rogen™ RNA UltraSense™ One-Step Quantitative RT- PCR System | | | | |
| | - | | - | | + | 13936 : 040 | QIAGEN QIAamp® Viral RNA QIAGEN OneStep RT-PCR kit | | | | |
| | - | | - | | + | Juna | BIOTECON Diagnostics Extraction Kit | | | | |
| | - | | - | | Not exal, inc. | | CONGEN SureFast® Prep DNA/RNA virus CONGEN SureFast® Norovirus PLUS | | | | |
| | | | | • | 0 | | BIOTECON Diagnostics Extraction Kit | | | | |

Questionnaire results:

Please note that not all participants provided the relevant information. The data analysed does not evaluate or associate the results with a failure with PT to a method/process used nor does it attempt to compare performance of the various molecular kits/processes with each other.

The graph below shows the participants that took part in this distribution from each country.



The RNA extractions used is shown in the table below:

| | No of users | % of users |
|--|-------------|------------|
| BioMérieux vucliS INS® miniMag® | 19 | 54 |
| BIOTECON i gnosti s Extraction Kit | 2 | 6 |
| CONGEN Sure Prep DNA/RNA virus | 4 | 11 |
| In-house | 1 | 3 |
| KingFisher™ MagMAX™ CORE Nucleic Acid Purification Kit | 1 | 3 |
| MACHEREY-NAGEL NucleoSpin® RNA Virus | 1 | 3 |
| QIAGEN QIAamp® Viral RNA | 3 | 9 |
| QIAGEN QIAsymphony DSP virus/pathogen mini kit | 1 | 3 |
| QIAGEN RNeasy kit | 1 | 3 |
| Roche High Pure Viral RNA kit | 1 | 3 |
| Stratec INVISORB Spin Virus RNA Mini Kit | 1 | 3 |

The PCR reagents used is shown in the table below:

| | No of users | % of users |
|---|-------------|------------|
| Altona DIAGNOSTICS RealStar® Norovirus RT-PCR Kit 2.0 | 1 | 3 |
| Applied Biosystems® TaqMan® Universal PCR Master Mix | 1 | 3 |
| Applied Biosystems™ AgPath-ID™ One-Step PCR Reagents | 1 | 3 |
| Applied Biosystems™ TaqMan™ Fast Virus One-Step Master Mix | 1 | 3 |
| CeeramTools® GI, GII and HAV detection kits | 9 | 28 |
| CONGEN SureFast® Norovirus PLUS | 1 | 3 |
| CONGEN SureFast® Norovirus/Hepatitis A 3plex kit | 2 | 6 |
| Invitrogen™ RNA UltraSense™ One-Step Quantitative RT-PCR System | 13 | 41 |
| ISO 15216-1:2017 MasterMix Invitrogen | 1 | 3 |
| QIAGEN OneStep RT-PCR kit | 1 | 3 |
| QIAGEN® QuantiTect® Probe RT-PCR kit | 1 | 3 |

End of report