

Summary of Results

External Quality Assessment of Water Microbiology Drinking Water Scheme

Distribution Number: W206 Sample Numbers: W206A, W206B

Distribution Date:	January 2023
Results Due:	
	10 February 2023
Report Date:	21 February 2023
Samples prepared and quality control tested by:	Divya George Nafeesa Hussain Cansev Katar Margaret Njenga Zak Prior Jake Videlefsky
Data analysed by:	Joanna Donn Nita Patel
Report compiled by:	Joanna Donn Nita Patel
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For further information on the scheme please refer to:

Scheme Guide: https://www.gov.uk/government/publications/food-and-water-proficiency-testing-schemes-scheme-guide

Guide to Scoring and Statistics:

https://www.gov.uk/government/publications/food-and-water-proficiency-testing-schemes-scoring-systems-and-statistics

General guidance for z-scores:

Participants' enumeration results are converted into z-scores using the following formula:

xi = participants' result expressed (expressed as a log 10 value)

 $Z = (X_1 - X_{pl})$ X_{pt} = assigned value (participants' consensus median (expressed as a log 10 value))

 σ_{pt} = the fixed standard deviation for the examination (calculated by FEPTU)

The σ_{pt} -value expresses the acceptable difference between the individual participant's result and the participants' consensus median. The σ_{pt} -value used for calculating z-scores for all parameters in the Drinking Water Scheme is 0.35. A guide to interpreting z-scores follows, although laboratories must interpret their scores in the context of their own laboratory situation.

z = -1.99 to +1.99 satisfactory z = -2 to -2.99 or +2 to +2.99 questionable z = < -3.00 or > +3.00 unsatisfactory

It is usually recommended that z-scores exceeding +/-2 are investigated to establish the possible cause. As a general rule, UKHSA recommends that all questionable and unsatisfactory results are investigated.

FEPTU Quality Control: To demonstrate homogeneity of the sample, a minimum of 10 LENTICULE® discs, selected randomly from a batch, are tested in duplicate for parameters requiring enumeration.

To demonstrate stability of the sample, a minimum of six LENTICULE discs, selected randomly from a batch, are examined throughout the distribution period for enumeration parameters.

UKHSA uses methods stipulated in the Environmental Agency Standing Committee of Analysts series of documents: Microbiology of Water and Associated Materials (2017) - Practices and Procedures for Laboratories. However for *Escherichia coli* and coliform examinations, ISO 9308-1:2014 Water quality - Enumeration of *Escherichia coli* and coliform bacteria Part 1: Membrane filtration method for waters with low bacterial background flora is followed.

The FEPTU results are used for guidance in the preliminary intended results notification, letters are posted on the website immediately after every distribution; electronic notification of their availability is sent to all participants.

Refer to section 17.0 of the Scheme Guide if you have experienced difficulties with any of the examinations.

https://www.gov.uk/government/publications/food-and-water-proficiency-testing-schemes-scheme-guide

Participants are reminded that reporting an incorrect or false negative results could have serious public health implications.

Results, as summarised in the performance assessment sheet included in this report, provide a more effective indication of on-going problems with the enumeration of low levels of indicator organisms in drinking water samples. Performance assessment cannot be undertaken as effectively if laboratories do not participate in all distributions for the Drinking Water Scheme.

Please contact FEPTU staff for advice and information:

Repeat samples Carmen Gomes or Kermin Daruwalla Tel: +44 (0)20 8327 7119

Data Analysis Nita Patel Fax:

Microbiological advice

Email: foodeqa@ukhsa.gov.uk

General comments and complaints

Nita Patel or Zak Prior

Scheme consultants Stephen Bullock

Scheme Co-ordinator Nita Patel

Accreditation: UKHSA Water EQA Scheme for Drinking Water is accredited by the United Kingdom Accreditation Service (UKAS) to ISO/IEC 17043:2010.



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Sample: W206A

Contents: Citrobacter freundii (14) (NCTC 9750), Pseudomonas aeruginosa (34) (wild strain), Clostridium perfringens (40) (wild strain),

Staphylococcus saprophyticus (29 per mL) (wild strain) - all levels are per 100mL unless otherwise stated

Expected Results:

All counts are expressed as colony forming units (cfu) per 100mL except the colony counts at 37° C/48h and 22° C/72h which are cfu per 100mL except the colony counts at 37° C/48h and 37° C/48h

nL.

mL.							
Parameter	Coliform bacteria	E.coli	Enterococci	P.aeruginosa	C.perfringens	Colony count 37°C/48h	Colony count 22°C/72h
FEPTU median	15	0	0	32	41	28	29
No. results returned	94	96	87	77	66	89	86
Assigned value (Participants median all results)	18	0	0	34	54	25	27
Uncertainty of assigned value*	0.07	N/A	N/A	0.04	0.02	0.02	0.02
Participants Algorithm A mean (all results)	18	0	0	33	52	25	26
Expected Range	1 - 57	N/A	N/A	11 - 106	17 - 171	8 - 79	8 - 84
Standard deviation** (log ₁₀)	0.49	N/A	N/A	0.24	0.11	0.14	0.12
No of outlying counts	16	N/A	N/A	10	5	5	5
False positives	N/A	0	5	N/A	N/A	N/A	N/A
False negatives	8	N/A	N/A	6	0	0	0
Your result							
Score for performance assessment							
Z-score							
Analyst 2 Result							
Analyst 2 Z-score							
Analyst 3 Result							
Analyst 3 Z-score							

^{*} U(Xpt) is based on results transformed to a log10 scale

^{**} Robust S^* based on median absolute deviation about the participants' median (MADe) and is based on logged data

Total sent samples	103
Not examined	2
Non returns	2
Late returns	0

Sample: W206B

Contents: Enterobacter cloacae (12) (wild strain), Escherichia coli (19) (wild strain), Enterococcus faecalis (53) (wild strain),

Pseudomonas aeruginosa (22) (wild strain), Clostridium perfringens (20) (wild strain), Microbacterium sp. (56 per mL) (wild

strain) - all levels are per 100mL unless otherwise stated

Expected Results:

All counts are expressed as colony forming units (cfu) per 100mL except the colony counts at 37°C/48h and 22°C/72h which are cfu per

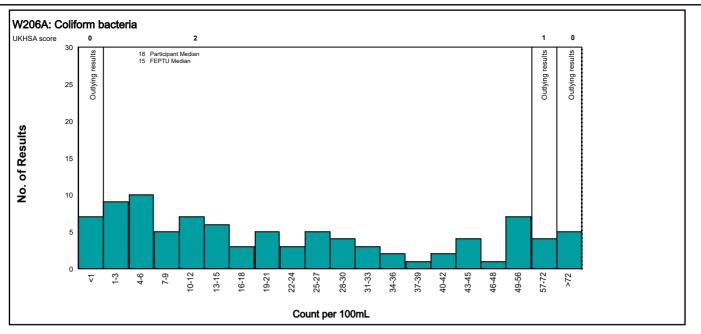
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IIIL.							
Parameter	Coliform bacteria	E.coli	Enterococci	P.aeruginosa	C.perfringens	Colony count 37°C/48h	Colony count 22°C/72h
FEPTU median	31	19	50	23	20	2	56
No. results returned	93	96	87	77	66	87	86
Assigned value (Participants median all results)	43	31	61	24	27	2	71
Uncertainty of assigned value*	0.03	0.03	0.01	0.03	0.03	0.04	0.02
Participants Algorithm A mean (all results)	41	30	61	22	27	2	70
Expected Range	14 - 136	10 - 96	19 - 193	8 - 76	9 - 85	0 - 6	22 - 225
Standard deviation** (log ₁₀)	0.21	0.23	0.1	0.23	0.16	0.25	0.12
No of outlying counts	3	3	2	8	5	3	4
False positives	N/A	N/A	N/A	N/A	N/A	N/A	N/A
False negatives	0	0	1	2	1	0	0
Your result							
Score for performance assessment							
Z-score							
Analyst 2 Result							
Analyst 2 Z-score							
Analyst 3 Result							
Analyst 3 Z-score							

^{*} $U(X_{pt})$ is based on results transformed to a log_{10} scale

^{**} Robust S^* based on median absolute deviation about the participants' median (MADe) and is based on logged data

Total sent samples	103
Not examined	3
Non returns	2
Late returns	0

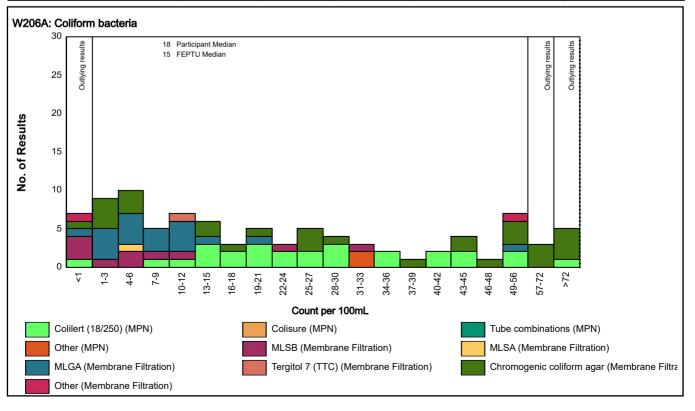


W206A: Coliform bacteria

FEPTU Method: Chromogenic coliform

agar

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range	e Rep	orted
Colilert (18/250)	27	0	29	25	0.24	0	-	84
Colisure	0	0	0					
Tube combinations	0	0	0					
Other (MPN)	2	0	2					
MLSB	10	0	10	5	0.31	0	-	32
MLSA	1	0	1					
MLGA	19	0	20	7	0.30	0	-	50
Tergitol 7 (TTC)	1	0	1					
Chromogenic coliform agar	30	1	32	29	0.53	0	-	194
Other (Membrane filtration)	2	0	2					



W206A: Escherichia coli						
		No data	ı for graph			
Method based presentati	on					
W206A : Escherichia coli			E E1	PTI I Method: C	hromogenic colifor	m
WZOOM . ESCIIOIICIIIA COII			aga			
Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S*	Range Reported

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range	Repo	orted
Colilert (18/250)	20	6	25	0	0.00	0	-	0
Colisure	0	0	0					
Tube combinations	0	0	0					
Other (MPN)	2	0	2					
TBX	0	0	0					
MLGA	14	5	18	0	0.00	0	-	0
MLSB	10	0	12	0	0.00	0	-	0
MLSA	1	0	1					
Tergitol 7 (TTC)	1	0	1					
Chromogenic coliform agar	28	6	36	0	0.00	0	-	0
Other (Membrane filtration)	1	0	1					

W206A: Escherichia coli			

No data for graph

/206A: Enterococci			_			
lethod based presentat	ion	No data	a for graph			
/206A : Enterococci			FE	PTU Method: S	+B	
Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S*	Range Reported
Enterolert (E/DW/250)	4	3	5		(=-0)	
Other (MPN)	0	0	0			
Enterococcus agar	6	1	8		1	

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No data for graph

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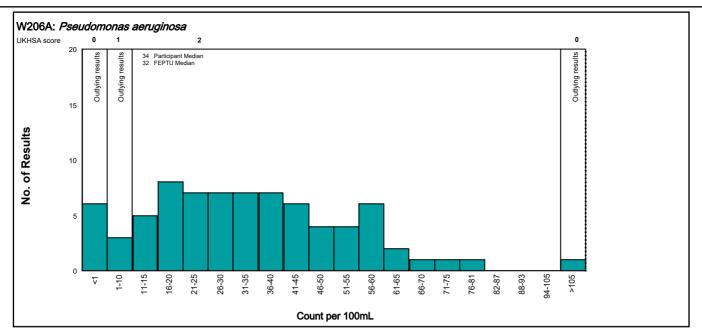
Other (Membrane filtration)

S+B

W206A: Enterococci

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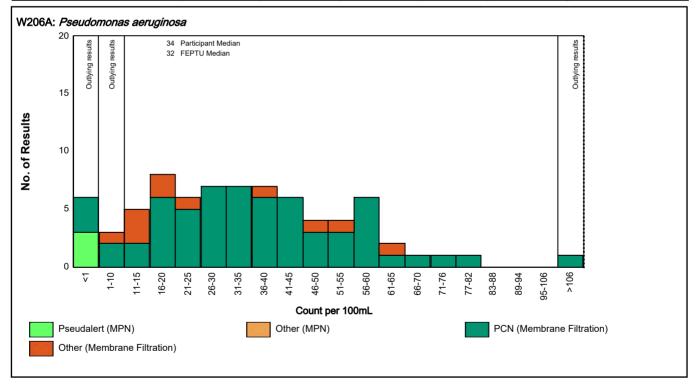
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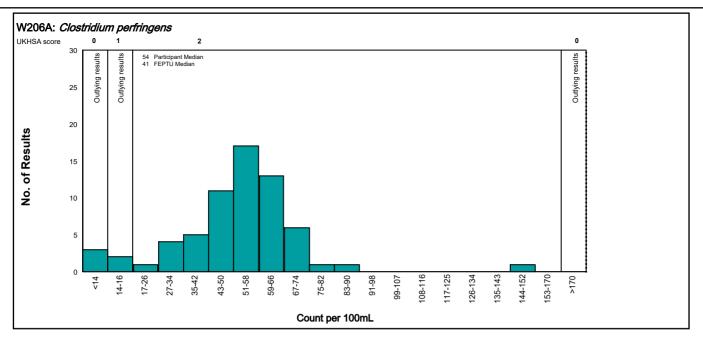


W206A: Pseudomonas aeruginosa

FEPTU Method: PCN

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Report	ed
Pseudalert	1	2	1				
Other (MPN)	0	0	0				
PCN	61	1	83	35	0.22	0 - 1	20
Other (Membrane filtration)	11	0	15	20	0.34	3 - 6	62

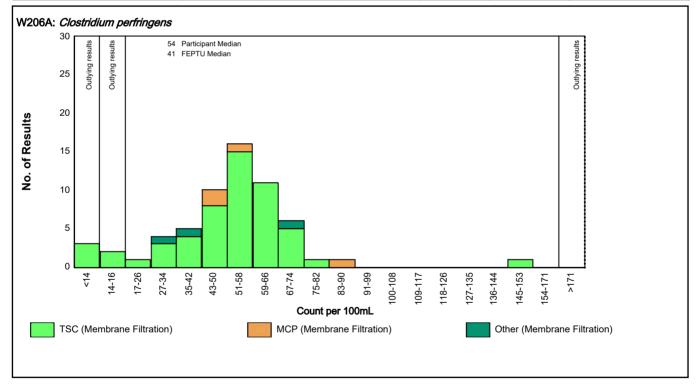


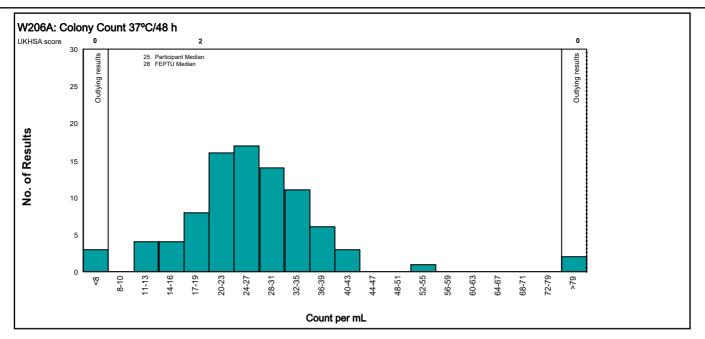


W206A: Clostridium perfringens

FEPT	U	Metho	d:	TSC
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Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
TSC	54	1	88	54	0.11	6 - 150
MCP	4	0	6			
Other (Membrane filtration)	3	0	4			

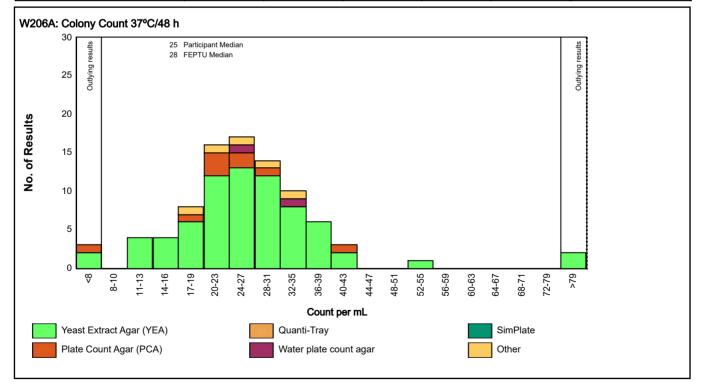


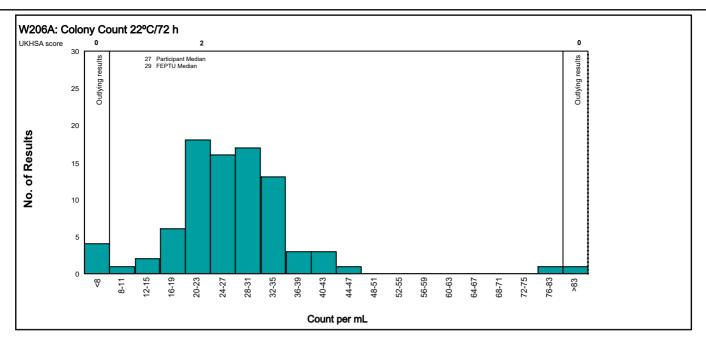


W206A: Colony Count 37°C/48 h

FEPTU Method: Yeast Extract Agar (YEA)

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
Yeast Extract Agar (YEA)	72	0	81	26	0.15	2 - 92
Quanti-Tray	0	0	0			
SimPlate	0	0	0			
Plate Count Agar (PCA)	9	0	10			
Water plate count agar	2	0	2			
Other	5	0	5			

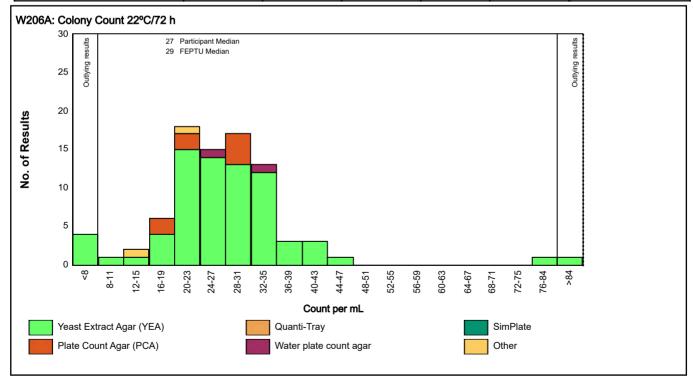


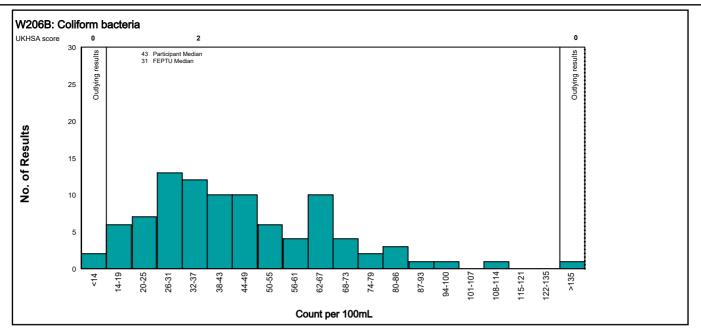


W206A: Colony Count 22°C/72 h

FEPTU Method: Yeast Extract Agar (YEA)

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Repo	rted
Yeast Extract Agar (YEA)	73	1	85	27	0.13	1 -	93
Quanti-Tray	0	0	0				
SimPlate	0	0	0				
Plate Count Agar (PCA)	8	0	9				
Water plate count agar	2	0	2				
Other	2	0	2				



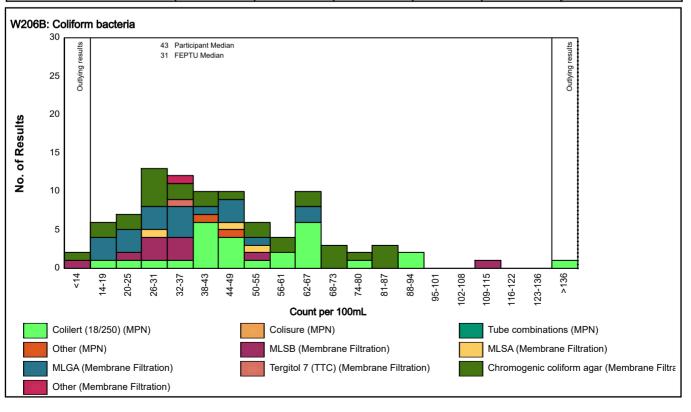


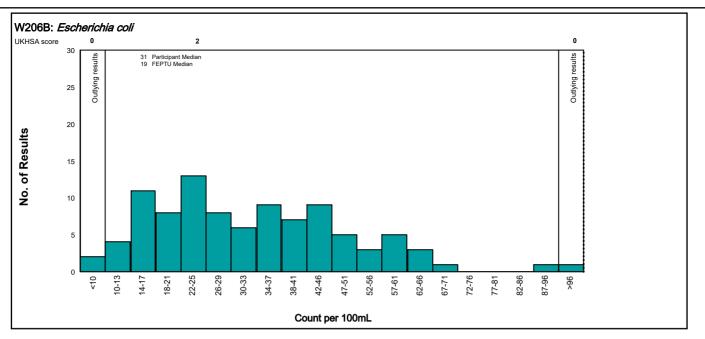
W206B : Coliform bacteria

FEPTU Method: Chromogenic coliform

agar

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range	Rep	orted
Colilert (18/250)	27	0	29	47	0.14	15	-	201
Colisure	0	0	0					
Tube combinations	0	0	0					
Other (MPN)	2	0	2					
MLSB	10	0	10	33	0.14	13	-	112
MLSA	3	0	3					
MLGA	20	0	21	34	0.21	15	-	66
Tergitol 7 (TTC)	1	0	1					
Chromogenic coliform agar	28	0	30	44	0.25	13	-	84
Other (Membrane filtration)	1	0	1					



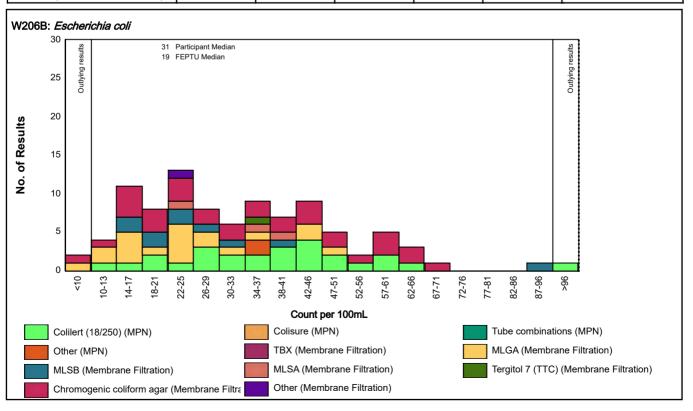


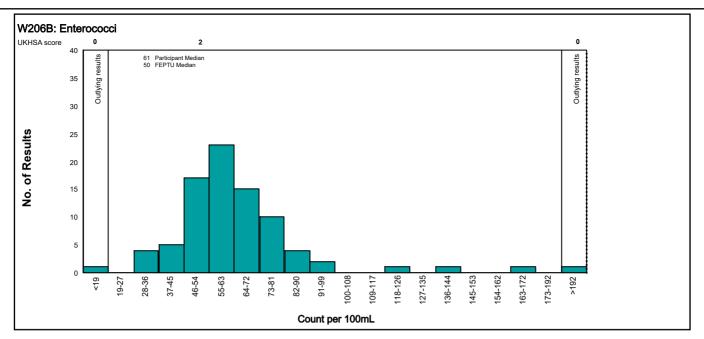
W206B: Escherichia coli

FEPTU Method: Chromogenic coliform

agar

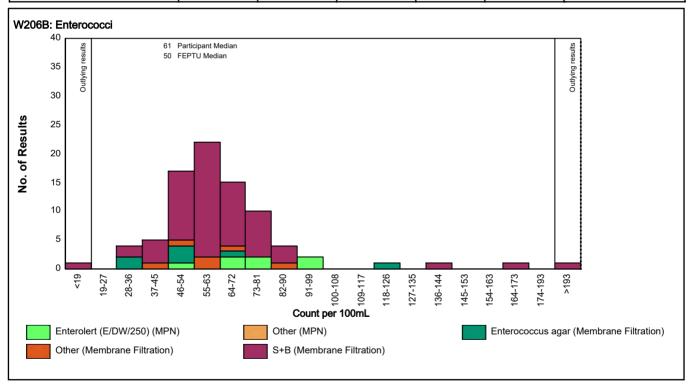
Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range	Rep	orted
Colilert (18/250)	26	0	27	38	0.19	12	-	182
Colisure	0	0	0					
Tube combinations	0	0	0					
Other (MPN)	2	0	2					
TBX	0	0	0					
MLGA	20	0	21	22	0.24	9	-	49
MLSB	10	0	10	23	0.16	16	-	95
MLSA	3	0	3					
Tergitol 7 (TTC)	1	0	1					
Chromogenic coliform agar	32	0	33	34	0.26	5	-	71
Other (Membrane filtration)	1	0	1					

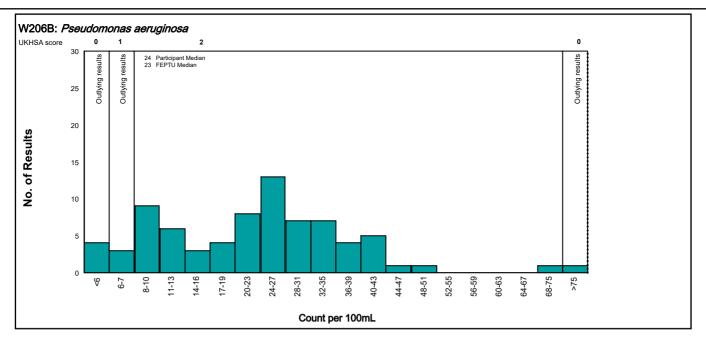




W206B : Enterococci FEPTU Method: S+B

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
Enterolert (E/DW/250)	7	0	8			
Other (MPN)	0	0	0			
Enterococcus agar	7	0	8			
Other (Membrane filtration)	6	0	7			
S+B	63	3	75	61	0.09	29 - 272

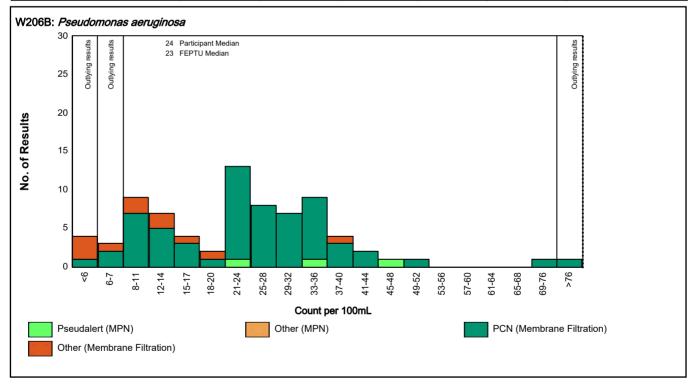


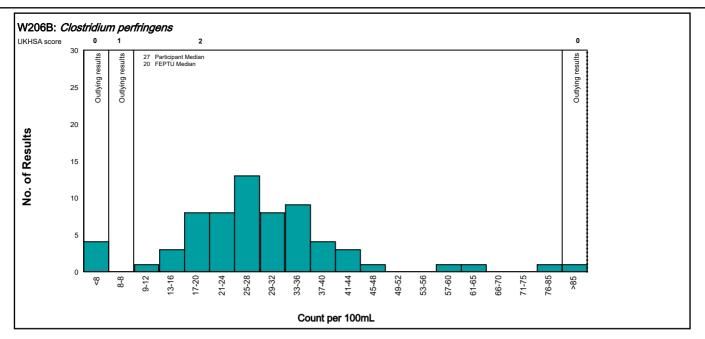


W206B: Pseudomonas aeruginosa

FEPTU Method: PCN

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
Pseudalert	3	0	3			
Other (MPN)	0	0	0			
PCN	62	0	81	25	0.22	4 - 80
Other (Membrane filtration)	11	0	14	9	0.26	0 - 38

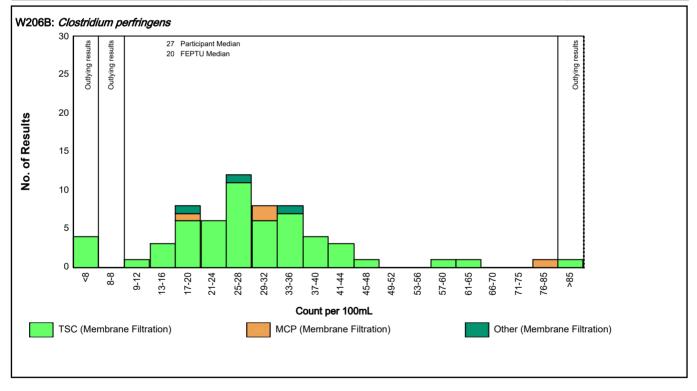




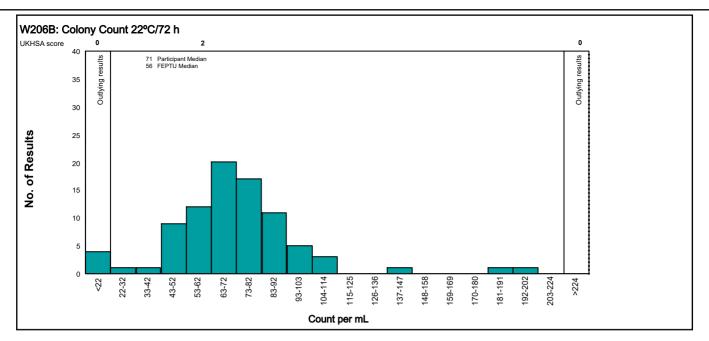
W206B: Clostridium perfringens

FEPT	J Method:	TSC
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Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
TSC	55	0	88	27	0.17	0 - 89
MCP	4	0	6			
Other (Membrane filtration)	3	0	4			



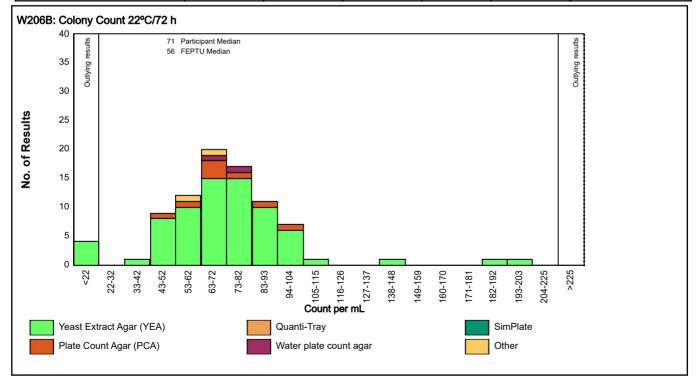
/206B: Colony Count 37°C/48 h							
		No data	a for graph				
		NO data	i loi grapii				
lethod based presentat	tion						
/206B : Colony Count 37°C/48 h			FEI (YE		east Extract Agar		
	N	Excluded	Percentage	Median	Robust S*	D	
			i Percentade i		I RANHET VAL		
Method	Number of Results			Wedian	1	Range	Reported
	Results	Results	of the total		(Log ₁₀)		
Yeast Extract Agar (YEA)				2	1	0	
	Results 69	Results 3	of the total		(Log ₁₀)		
Yeast Extract Agar (YEA) Quanti-Tray SimPlate	Results 69 0	Results 3 0	of the total 83		(Log ₁₀)		
Yeast Extract Agar (YEA) Quanti-Tray	69 0	3 0 0	of the total 83 0		(Log ₁₀)		



W206B: Colony Count 22°C/72 h

FEPTU Method: Yeast Extract Agar (YEA)

Method	Number of Results	Excluded Results	Percentage of the total	Median	Robust S* (Log ₁₀)	Range Reported
Yeast Extract Agar (YEA)	73	1	85	72	0.12	2 - 202
Quanti-Tray	0	0	0			
SimPlate	0	0	0			
Plate Count Agar (PCA)	8	0	9			
Water plate count agar	2	0	2			
Other	2	0	2			



Performance Assessment Sheet

Distribution	Sample	Coliform bacteria score	Escherichia coli score	Enterococci score	Pseudomonas aeruginosa	Clostridium perfringens	Colony Count 37°C/48 h	Colony Count 22°C/72 h
	W206A				sone	Strie	State	SCHE
W206	W206B							
	W205A							
W205	W205B							
	W204A							
W204	W204B							
	W203A							
W203	W203B							
	W202A							
W202	W202B							
	W201A							
W201	W201B							
Total max possible								
Total perc	entage							

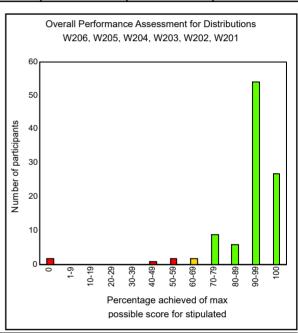
Performance Assessment Comment:

Participants are reminded that to take advantage of the performance assessment overtime tool provided in the reports they need to take part in more than one distribution a year.

Performance assessments are designed to alert participants to on-going problems with their examinations and are provided after every distribution. Scores are allocated to results reported for every parameter, for every sample to help assess performance.

Cummulative scores are calculated for the current and previous **five** distributions for the Drinking Water Scheme. Participants' cummulative scores for each of the examinations are compared with the maximum possible scores after every distribution.

Your overall performance with the enumerations of low levels of indicator organisms in drinking water proficiency testing samples for the current and last five distributions is collated in the chart to the right.



Performance Assessment Comment:

Laboratories that achieve less than 70% of the maximum possible score are likely to be experiencing significant problems with their examinations and are advised to:

- a) refer to the relevant distribution reports for sample-specific comments
- b) refer to the website guidance documents:
 https://www.gov.uk/government/collections/external-quality-assessment-eqa-and-proficiency-testing-pt-for-food-water-and-environmental-microbiology

Sample specific comment

W206A: Coliform

This sample contained a Citrobacter freundii which is a Gram negative coliform bacteria in the Enterobacteriaceae family. In the FEPTU laboratory this organism grew as <0.5mm pink translucent colonies on coliform chromogenic agar following incubation at 37°C for 24 hours.

8/94 (9%) laboratories reported a false negative result for this examination, the standard deviation was also wide. A Kolmogorov–Smirnov (KS) test showed that the data was evenly distributed. Laboratories reporting an incorrect PT result should request a repeat sample for their own internal investigation.

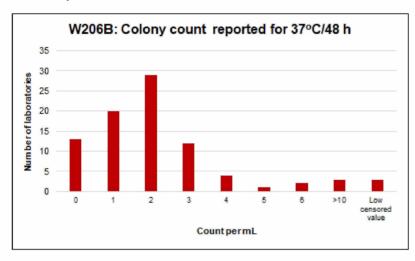
Please note that the species of the *Citrobacter* was incorrectly shown on the intended results published on the 13 February 2023. This sample contained a *C. freundii* and not *C. braakii*, this error will not have an impact on the results reported and the performance of the laboratories with this examination. FEPTU's reference number is FNC742.

W206A: Pseudomonas aeruginosa

This strain of *P. aeruginosa* generates a negative reaction in the Pseudalert method. This suggests that this strain does not have the enzyme that cleaves the substrate in the reagent to produce a blue fluorescence under ultraviolet light for results to be positive in the Pseudalert system. Participants reporting a false negative result that used a Pseudalert have been score correctly. It is important that participants understand the limitations of the methods they use and the impact this may have on public health.

W206B: Colony count 37°C/48 h

Below is a graph of the colony counts reported at 37°C/48h by the participants, which is not shown in the main scheme report.



General comments:

Method based presentation of results:

Participants are advised if less than 10 laboratories report a result for a method, no data is shown for the Median, Robust S* and the Range Reported in the 'Method based presentation' tables.

Numbers shown in the 'Excluded Results' column are laboratories that reported a censored value.

Trend analysis:

Plotting your PT results over a period can help to identify potential problems. If you need the latest file, please email us on footega@ukhsa.gov.uk.

If you do not return a result for a distribution, you will not be able to view all the participants' results data in your individualised report. Please contact us if you require this information on footega@ukhsa.gov.uk.

End of report

