

# **QMS**

### **Quality in Microbiology Scheme**

## **Scheme Description**

# LGC Proficiency Testing

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#### Record of issue status and modifications

ISSUE	ISSUE DATE	DETAILS	AUTHORISED BY
20	Sep 2020	Added samples for detection of Listeria in vegetables and cheese.  Added sample 45 thermophilic acidophilic bacteria in fruit.	L. Chesters
21	Sep 2020	Added additional samples in herbs and spice matrix	T Noblett
22	July 2021	Reformatted numbering for Salmonella by changing 28 to 06T, 30 to 06H and 31 to 06S. Added additional samples 06CF and 46EG Updated email address and UKAS logo	T.Noblett
			A Collins
23	Feb 2022	Added combined analyte for sample 27. Added Enterobacteriaceae analyte for sample 41.	C Taylor
24	Sept 2022	Added A and B to sample 46EG to show 2 different samples within the package.  C h a n g e d t h e f o r mat oto state seachassatiyte a n d separately for any relevant analytes.  Added combined osmophilic yeast and osmophilic mould analyte to sample 05 and changed format to state each analyte separately. Sample range updated to 0 – 100 cfu/g for sample 04.  Addition of samples 47 (Detection of Listeria species in pooled sample), 48 (Pathogens in Infant formula), 49 (Microscopical Identification) and 50 (Indicator organisms in cheese).	C Taylor
25	Nov 2022	Cronobacter in 10g added to sample 48 and corrected reporting units	A Cheetham
26	Apr 2023	Added E.coli as analyte to sample 41.	M.Bell
27	Sept 2023	Range updated for sample 27.  Samples 06 EG,CF and VG updated to vial and matrix. Sample 21 changed to vial and >25g matrix.  Ne w a n a l y t e a d d e dCarhpsylobacterDsept eecci tei so'n cfu/g. Sample 32 changed to vial and 10g matrix.  New samples 51, 52, 53, 54 added  A d d e d -t'onxoing e n i oE.codi O15a samples t o Removed sample 49  A d d ' l d e n t i f i c a t i o n o f S a l monel l A d d ' l d e n t i f i c a t i o n o f Y e a s t a n d S a mp l e s 5 2, 5 3 a n d 5 4 marked a s o f L G C' s f l e x i b l e s c o p e o f UKAS	M.Bell T.Noblett N. Mason
28	Mar 2024	Updated the descriptions of sample 52 and 53	A Collins
29	Apr 2024	Sample 51 'supplied as' updated	M.Bell
30	July 2024	Sample 55 added. Sample 40, 43 and 47 format updated to ready to test.	M.Bell
31	Oct 2024	Matrices removed from samples 21 and 32. Range for sample 47 amended to 0 to 100 cfu/125g. Format change for sample 47 from RTT to vial and matrix.	M.Bell
32	Feb 2025	Added new sample PT-MC-56 (A & B) for Salmonella in Therapeutic foods.	T.Noblett

#### Notes:

Where this document has been translated, the English version shall remain the definitive version.

#### **Scheme Aims and Organisation**

The primary aim of the Quality In Microbiology (QMS) is to enable laboratories performing the microbiological analysis of food and dairy products to monitor their performance and compare it with that of their peers. QMS also aims to provide information to participants on technical issues and methodologies relating to testing of food and dairy products.

The QMS scheme year operates from January to December. Further information about QMS, including test material availability, round despatch dates and reporting deadlines, are available on the current QMS application form.

#### **Test Materials**

Details of test materials available in QMS are given in Appendix A. The test parameters are continually reviewed to ensure they meet the needs of current laboratory testing and regulatory requirements.

Test material batches are tested for homogeneity for at least one test parameter where deemed appropriate. Details of homogeneity tests performed and results are given in the QMS Scheme Reports.

Some aspects of the scheme, such as test material production, homogeneity testing and stability assessment, can from time to time be subcontracted. When subcontracting occurs, it is placed with a competent subcontractor and LGC is responsible for this work. The planning of the scheme, the evaluation of performance and the authorisation of the final report will never be subcontracted.

#### **Statistical Analysis**

Information on the statistics used in QMS can be found in the General Protocol and in the Scheme Report. Methods for determining assigned values and the values for SDPA used for individual samples are given in Appendix A

#### **Methods**

Methods are listed in PORTAL. Please select the most appropriate method from the list. If none of the methods are appropriate, then please description in the Comments Section in PORTAL.

#### **Results and Reports**

QMS results are returned through our electronic reporting software, PORTAL, full instructions for which are provided by email.

QMS reports will be available on the website within 10 working days of round closure. Participants will be emailed a link to the report when it is available.

#### **APPENDIX A - Description of abbreviations used**

#### Assigned Value (AV)

The assigned value may be derived in the following ways:

§ From the robust mean (RMean). This is the median of participant results after the removal of test results that are inappropriate for statistical evaluation, e.g. miscalculations, transpositions and other gross errors. Generally, the assigned value will be set using results from all methods, unless the measurement is considered method-dependant, in which case the assigned value will be set by method as illustrated in the report tables.

For some analytes, where there is a recognised reference method for that type of measurement, this may be used as the assigned value for a particular analyte i.e. it would be applied to results obtained by any method.

Traceability: Assigned values which are derived from the participant results, or a sub-set of the results are not traceable to an international measurement standard. The uncertainty of assigned values derived in this way is estimated from the participant results, according to ISO 13528.

š From a formulation value (Formulation). This denotes the use of an assigned value derived from sample preparation details, where known and exact quantities of analyte have been used to prepare the sample.

Traceability: Assigned values calculated from the formulation of the test sample are traceable, via an unbroken metrological traceability chain, to an international measurement standard. The measurement uncertainty of the assigned value is calculated using the contributions from each calibration in the traceability chain.

š From a qualitative formulation (Qual Form). This applies to qualitative tests where the assigned value is simply based on the presence/absence of the analyte in the test material.

Traceability: Assigned values calculated from the qualitative formulation of the test sample are traceable to a certified reference standard or a microbiological reference strain.

š From expert labs (Expert). The assigned laboratory.

V¦ æ&^ æà ã | ã c ^ K Á Œ• • ã \* } ^ å Á ç æ| ˇ ^ • Á ] ¦ [ ç ã å ^ å Á à international measurement standard, according to the laboratory and the method used. The uncertainty of measurement for an assigned value produced in this way will be provided by the laboratory undertaking the analysis. Details of traceability and the associated uncertainty will be provided in the report for the scheme/round.

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#### Range

This indicates the concentration range at which the analyte may be present in the test material.

#### **SDPA**

The SDPA represents the 'standard deviation of assess participant performance for the measurement of each analyte. This may be a fixed value (as stated), a percentage (%) of the assigned value or based on the robust standard deviation of the participant measurement results, either across all methods or by method depending on whether the measurement made is method dependent (see assigned value).

#### Units

This indicates the units used for the assessment of data and in which participants should report their results. For some analytes in some schemes participants may have a choice of which units to report their results, however, the units stipulated in this scheme description are the default units to which any results reported using allowable alternative results will be converted to.

#### DP

This indicates the number of decimal places to which participants should report their measurement results.

#### **APPENDIX A**

Sample PT-MC-03 Enumeration of Salmonella species

**Supplied as:** 1 x 10g skimmed milk powder

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of Salmonella species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-04 Detection of Cronobacter species
Supplied as: 1 x 25g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of <i>Cronobacter</i> species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of <i>Cronobacter</i> species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA

Sample PT-MC-05 Enumeration of osmophilic yeast and mould (ISO 21527-2)

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of osmophilic yeast	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of osmophilic mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of osmophilic yeast and osmophilic mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-06 Supplied as: **Detection of Salmonella species** 

**06F** – Salmonella in generic food product - 25g oatmeal

**06D** – Salmonella in dairy/milk product - 25g skimmed milk powder **06EG** – Salmonella in egg products - 25g egg powder + 10ml vial

06CF - Salmonella in chicken faeces - 25g dried chicken faeces + 10ml vial

**06VG E** Salmonella in salads and vegetables - 25g dried mixed vegetables + 10ml vial

06CH - Salmonella in cheese - 25g cheese + 10ml vial

06NS E Salmonella in seeds and nuts - 25g seeds and/or nuts + 10ml vial

**06TE** Ë Salmonella in tea – 25g tea + 10ml vial

**06HB E** Salmonella in herbs – 25g dried herb + 10ml vial **06SP E** Salmonella in spices – 25g pepper or spice + 10ml vial

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Salmonella species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-07 Supplied as:

**Detection of Listeria species** 

07F - Listeria in food - 25g oatmeal

**07D** – Listeria in dairy/milk - 25g skimmed milk powder

**07VG** – Listeria in salads and vegetables - 25g dried mixed vegetables

**07CH** – Listeria in cheese - 25g cheese + 10ml vial **07HB** – Listeria in herbs - 25g herb + 10ml vial

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Listeria species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Listeria monocytogenes	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-08

**Enumeration of Listeria species** 

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of <i>Listeria</i> species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Listeria monocytogenes	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-09

**Enumeration of Enterococci** 

Supplied as:

1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of Enterococci	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-10 Enumeration of Clostridium species
Supplied as: 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Clostridium species	ALL	Qual Form	0 to 100,000	NA	Detected/Not detected 10g	NA
Detection of Clostridium perfringens	ALL	Qual Form	0 to 100,000	NA	Detected/Not detected 10g	NA
Enumeration of Clostridium perfringens	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Clostridium species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-11 Spore counts

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of mesophilic aerobic spores	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0
Thermophilic aerobic plate count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of thermophilic aerobic spores	Heated for 10 min @ 80°C Heated for 12 min @ 100°C	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0
Enumeration of highly heat resistant thermophilic aerobic spores	Heated for 30 min @ 100°C	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0

Sample PT-MC-12 **Detection of Shigella species** 

Supplied as: 1 x 25g oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Shigella species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-13 Supplied as: **Detection of Vibrio species** 

1 x 25g oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Vibrio species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Vibrio parahaemolyticus	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-14 **Detection of Yersinia species** Supplied as: 1 x 25g skimmed milk powder

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Yersinia species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Yersinia enterocolitica	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-15 Anaerobes

Supplied as: 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total anaerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of anaerobic sulphite-reducing bacteria	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of mesophilic anaerobic spores	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0
Enumeration of sulphite-reducing <i>Clostridium</i> spores	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0

Sample PT-MC-16

TVC/indicator organisms
1 x 10g skimmed milk powder or oatmeal Supplied as:

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Enterobacteriaceae	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Escherichia coli	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-17 Enumeration of Staphylococcus and Bacillus species

Supplied as: 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of Staphylococcus species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coagulase-positive Staphylococci	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Bacillus species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Bacillus cereus	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-18 Low-level indicator organisms

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Escherichia coli	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA
Detection of Enterobacteriaceae	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA
Detection of coliforms	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA
Enumeration of Escherichia coli	ALL	RMean	0 to 1,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Enterobacteriaceae	ALL	RMean	0 to 1,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 1,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-20 Enumeration of thermotolerant coliforms
Supplied as: 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of Escherichia coli	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of thermotolerant coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-21 Detection of Campylobacter species
Supplied as: 1 x 10ml vial to represent 10g or 25g sample

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Campylobacter species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Campylobacter species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA

Sample PT-MC-22 Detection of E.coli O157 (non-toxigenic strain)

**Supplied as:** 1 x 25g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Escherichia coli O157	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-23 Enumeration of yeast and mould (ISO 21527-1)

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of yeast	ALL	RMean	0 to 100,000	log10 0.35	cfu/g	0
Enumeration of mould	ALL	RMean	0 to 100,000	log10 0.35	cfu/g	0
Enumeration of yeast and mould combined	ALL	RMean	0 to 100,000	log10 0.35	cfu/g	0
Identification of yeast and mould to genus and/or species level	ALL	Formulation	NA	NA	NA	NA

Sample PT-MC-24 Supplied as:

**Enumeration of Lactic acid bacteria** 

024F - Lactic acid bacteria in food - 10g oatmeal

**024D** – Lactic acid bacteria in dairy/milk - 10g skimmed milk powder **024HB** – Lactic acid bacteria in herb - 10g dried herb + 10ml vial **024SP** – Lactic acid bacteria in spice- 10g pepper or spice + 10ml vial

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of lactic acid bacteria	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-25 Psychrotrophs

**Supplied as:** 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of aerobic psychrotrophs	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-26 Pseudomonas species

**Supplied as:** 1 x 10g skimmed milk powder or oatmeal

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of <i>Pseudomonas</i> species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Detection of Pseudomonas species	ALL	Qual Form	0 to 100,000	NA	Detected/Not detected 10g	NA
					-	

Sample PT-MC-27 Enumeration of probiotic bacteria

**Supplied as:** 1 x 10ml vial to represent 10g sample (once reconstituted in 10ml diluent)

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of <i>Bifidobacterium</i> species	ALL	RMean	0 to 10,000,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Lactobacillus species	ALL	RMean	0 to 10,000,000	log <sub>10</sub> 0.35	cfu/g	0
Combined enumeration of Lactobacillus species and Bifidobacterium species	ALL	RMean	0 to 20,000,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-29 Indicator organisms in tea
Supplied as: 1 x 10ml vial plus 10g tea matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coagulase positive Staphylococci	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of yeast	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of yeast and mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-32 Enumeration of Campylobacter species

**Supplied as:** 1 x 10ml vial to represent 10g sample (once reconstituted in 10ml diluent)

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of Campylobacter species	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.50	cfu/g	0

Sample PT-MC-33 Identification Test (non-pathogen)

**Supplied as:** 1 x 10ml vial containing a single organism. The sample may contain biosafety level 1 or 2 organisms, including Staphylococcus, Bacillus and Clostridium, but will not contain the recognised food pathogens

such as Salmonella, Listeria, Campylobacter or STEC.

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
	ALL	Formulation	NA	NA	NA	NA
genus or species level.						

Sample PT-MC-34 Salmonella identification

Supplied as: 1 x 10ml vial containing a single organism

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Salmonella identification to serogroup or serovar	ALL	Formulation	NA	NA	NA	NA

Sample PT-MC-35 Paper exercise

**Supplied as:** Participants will be provided with a photograph and a scenario in order to count the number of colonies

and calculate the number of microorganisms in the original sample.

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Counting of colonies and calculation of	Visual count only	Formulation	NA	Greater of robust	cfu/g	NA
number of microorganisms				SD or log 0.05		

Sample PT-MC-36 (A & B)
Supplied as:
Quantitative Package
36D EQuantitative in da

**36D E**Quantitative in dairy/milk - 2 x 10ml vial plus minimum 20g skimmed milk powder matrix

**36HB** – Quantitative in herbs – 2 x 10ml vial plus minimum 20g herb matrix

**36SP** – Quantitative in spice – 2 x 10ml vial plus minimum 20g pepper or spice matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Enterobacteriaceae	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Escherichia coli	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Bacillus cereus	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coagulase-positive	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Staphylococci						
Detection of coagulase-positive	ALL	QualForm	0 to 100,000	NA	Detected/Not detected	NA
Staphylococci					10g	
Enumeration of yeast	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of yeast and mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-37 (A & B) Qualitative Package

**Supplied as:** 37D - 2 x 10ml vial plus minimum 200g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Escherichia coli O157	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected	NA
(non-toxigenic strain)					25g	
Detection of <i>Listeria</i> species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected	NA
					25g	
Detection of Listeria	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected	NA
monocytogenes					25	
Identification of Listeria species	ALL	Formulation	NA	NA	NA	NA
Detection of Salmonella species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected	NA
					25g	
Identification of Salmonella	ALL	Formulation	NA	NA	NA	NA
species						

Sample PT-MC-38 Detection of Clostridium species and Staphylococcus species in milk

**Supplied as:** 1 x 10ml vial plus minimum 20g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Clostridium perfringens	ALL	Qual Form	0 to 100	NA	Detected/Not detected 10g	NA
Detection of coagulase positive Staphylococci	ALL	Qual Form	0 to 100	NA	Detected/Not detected 10g	NA

Sample PT-MC-39 Bacterial count in milk

**Supplied as:** 1 x 10g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/ml	SDPA	Reporting units	DP
Bacterial level by Bactoscan	Bactoscan	RMean	ALL	TBC	Bacteria/ml	NA
Bacterial level by colony count	ALL	RMean	ALL	TBC	cfu/ml	NA

Sample PT-MC-40 Detection of Salmonella species in pooled sample

**Supplied as:** 375g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/375g	SDPA	Reporting units	DP
Detection of Salmonella species	ALL	Qual Form	0 to 100	NA	Detected/Not detected 375g	NA

Sample PT-MC-41 Supplied as:

= b X ] WUhcf 'UbX'gdc] `U[Y'cf[Ub]gag']b' ÏfYUXm'aYU`Đ 1 x 10ml vial plus 1 x 10g dried food matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Escherichia coli	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Enterobacteriaceae	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coagulase positive Staphylococci	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of yeast	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of yeast and mould	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

Sample PT-MC-42 Detection of Cronobacter species at low-level

Supplied as: 1 x 10ml vial plus 1 x 25g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/25g	SDPA	Reporting units	DP
Detection of Cronobacter species	ALL	Qual Form	0 to 50	NA	Detected/Not detected 25g	NA

Sample PT-MC-43 Detection of Cronobacter species in 375g

**Supplied as:** 375g skimmed milk powder matrix

Analyte	Method	AV	Range cfu/375g	SDPA	Reporting units	DP
Detection of Cronobacter species	ALL	Qual Form	0 to 50	NA	Detected/Not detected 375g	NA

Sample PT-MC-44 Supplied as: DUh\c[Ybg'-fp-bYUhfDYUkmX

1 x 10ml vial plus minimum 100g dried processed food e.g. crisps, cakes, biscuits, confectionary, cereal

and other snacks.

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of <i>E. coli</i> O157 (non-toxigenic	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
strain)						
Detection of Listeria species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Listeria monocytogenes	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Identification of Listeria species	ALL	Formulation	NA	NA	NA	NA
Detection of Salmonella species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-45
Supplied as:

Thermophilic acidophilic bacteria
1 x 10ml vial plus 10g fruit matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Enumeration of thermophilic	MF	RMean	0 to 1,000	log <sub>10</sub> 0.35	cfu/g	0
acidophilic bacteria	BAT AGAR					
Detection of guaiacol producing	MF	Qual Form	0 to 1,000	NA	Detected/Not detected 10g	NA
thermophilic acidophilic bacteria	BAT AGAR		·			

Sample PT-MC-46 (A & B) Detection of Salmonella Typhimurium and/or Enteritidis in Egg powder

**Supplied as:** 46EG – Egg products – 2 x 25g egg powder

Analyte	Method	AV	Range cfu/g	SDPA	Reporting Units	DP
Detection of Salmonella Typhimurium	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Salmonella Enteritidis	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA

Sample PT-MC-47 Detection of Listeria species in 125g milk powder

**Supplied as:** 1 x 10ml vial plus 125g skimmed milk powder

Analyte	Method	AV	Range cfu/125g	SDPA	Reporting Units	DP
Detection of <i>Listeria</i> species	ALL	Qual Form	0 to 100	NA	Detected/Not detected 125g	NA
Detection of Listeria monocytogenes	ALL	Qual Form	0 to 100	NA	Detected/Not detected 125g	NA

Sample PT-MC-48 (A & B)\*

Pathogens in Infant formula

Supplied as:

2 x 10ml vials with minimum 100g infant formula

Analyte	Method	AV	Range cfu/g	SDPA	Reporting Units	DP
Detection of Salmonella species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Cronobacter species	ALL	Qual Form	0 to 1,000	NA	Detected/Not detected 25g	NA
Detection of Cronobacter species	ALL	Qual Form	0 to 2,500	NA	Detected/Not detected 10g	NA

<sup>\*</sup>Currently not included in LGC's UKAS Scope of Accreditation

Sample PT-MC-50\* Indicator organisms in cheese

**Supplied as:** 1 x lyophilised 10ml vial plus 10g dried cheese matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of Enterobacteriaceae	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0
Enumeration of coagulase-positive Staphylococci	ALL	RMean	0 to 100,000	log <sub>10</sub> 0.35	cfu/g	0

<sup>\*</sup>Currently not included in LGC's UKAS Scope of Accreditation

Sample PT-MC-51\* Commercial Sterility testing in milk

**Supplied as:** 5 x 10ml Lyophilised milk in vials which may or may not contain microorganisms at low levels.

Analyte	Method	AV	Range cfu/vial	SDPA	Reporting Units	DP
Commercial sterility	ALL	Qual Form	<100	N/A	Sterility Pass/Sterility	N/A
					Fail	
Identification of microorganisms	ALL	Formulation	NA	NA	NA	NA

<sup>\*</sup>Currently not included in LGC's UKAS Scope of Accreditation

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Sample PT-MC-52<sup>‡</sup> Gram-

**Gram-negative panel** 

Supplied as: 3 x 10ml vials each containing single organism (contents may include

pathogens and/or non-pathogens)

Analyte	Method	AV	Range cfu/vial	SDPA	Reporting Units	DP
Confirmation and/or identification of Gram-negative organisms (including	ALL	Formulation	NA	N/A	NA	N/A
Salmonella) to genus and/or species						

Sample PT-MC-53<sup>‡</sup> Gram-positive panel

**Supplied as:** 3 x 10ml vials each containing single organism (contents may include

pathogens and/or non-pathogens)

Analyte	Method	AV	Range cfu/vial	SDPA	Reporting Units	DP
Confirmation and/or identification of	ALL	Formulation	NA	N/A	NA	N/A
Gram-positive organisms (including						
Listeria) to genus and/or species						

Sample PT-MC-54<sup>‡</sup> Mixed culture for identification

**Supplied as:** 1 x 10ml vial containing up to 5 different organisms (contents may include

pathogens and/or non-pathogens)

Analyte	Method	AV	Range cfu/vial	SDPA	Reporting Units	DP
Identification of organisms to genus	ALL	Formulation	NA	N/A	NA	N/A
and/or species						

<sup>&</sup>lt;sup>‡</sup>Accredited within the boundaries of LGC's flexible scope of UKAS accreditation

Sample PT-MC-55\* Indicator organisms in nuts

**Supplied as:** 1 x lyophilised 10ml vial plus 10g nut matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting Units	DP
Total aerobic mesophilic count	ALL	RMean	0 to 100,000	0.35	cfu/g	0
Enumeration of coliforms	ALL	RMean	0 to 100,000	0.35	cfu/g	0
Enumeration of yeast	ALL	RMean	0 to 100,000	0.35	cfu/g	0
Enumeration of mould	ALL	RMean	0 to 100,000	0.35	cfu/g	0
Enumeration of yeast and mould	ALL	RMean	0 to 100,000	0.35	cfu/g	0

Sample PT-MC-56 (A & B)\* Detection of Salmonella species in foods for malnutrition of therapeutic use.

**Supplied as:** 2 x 25g high lipid and protein peanut paste matrix

Analyte	Method	AV	Range cfu/g	SDPA	Reporting units	DP
Detection of Salmonella species	ALL	Qual Form	0 to 1000	NA	Detected/Not detected 25g	NA

<sup>\*</sup>Currently not included in LGC's UKAS Scope of Accreditation

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