

Meat & Fish (QMAS) MT4871 - CLG Chemisches Labor Dr. Graser Individual Report

Round: 266

Issue Number 1 Issued 26 July 2018





Sample Details

Samples were despatched on 25 June 2018 The reporting deadline was 20 July 2018 The following samples were distributed in QMAS Round 266:

| Sample 735 | Matrix 10g Lyophilised meat | Contents Escherichia coli Klebsiella oxytoca |
|----------------------|---|---|
| 736 | 25g Lyophilised meat | Salmonella Senftenberg |
| 737 | 25g Lyophilised meat | Listeria monocytogenes |
| 738 | 10g Lyophilised meat | Staphylococcus epidermidis Clostridium perfringens |
| 746 | Lyophilised vial with 10g lyophilised meat | Pseudomonas aeruginosa Lactobacillus casei Candida albicans |
| 756A | Lyophilised vial with 10g meat powder | Enterobacter aerogenes Bacillus cereus Staphylococcus aureus Penicillium chrysogenum Saccharomyces cerevisiae |
| 756B | Lyophilised vial with 10g meat powder | Enterobacter aerogenes Escherichia coli Staphylococcus aureus Penicillium chrysogenum |
| 757A | Lyophilised vial with 100g meat powder | Escherichia coli O157 (non toxigenic strain) Listeria welshimeri Salmonella Indiana |
| 757B | Lyophilised vial with 100g meat powder | Escherichia coli O157 (non toxigenic strain) Listeria monocytogenes Listeria welshimeri |

Further information regarding assigned values, performance assessment and technical comments can be found under the individual sample and analyte results.

Individual Report

This individual report contains a summary of all the results submitted and the performance assessments for your laboratory and your individual analysts. Please note that nominated laboratory results are represented by a blue highlight in the analyst box.

Data statistics given in the individual report are for the method you have used for each analyte. Further detail can be obtained from the main report.

Full details of the scheme, sample types, analytes and data analysis can be found in the corresponding Main Report, along with any technical comments, if applicable. The Main Report is the definitive version.

If you have any questions regarding your results which are not answered in the Main Report, please contact us using the details on the front of the report. If you would like to order any samples for re-test, please contact our customer service department or your local office.

Results Summary

| Sample | Recults Reported | | | Unsatisfactory Results | Not Assessed^ |
|--|------------------|---|---|---------------------------|---------------|
| 735 - Meat Indicator Combination | 4 | 4 | 0 | 0 | 0 |
| 736 - Meat Salmonella (P/A) | 1 | 1 | 0 | 0 | 0 |
| 738 - Meat Clost/Staph | 1 | 1 | 0 | 0 | 0 |
| 746 - Pseudomonas/Lactic Acid/Yeast/Mould | 3 | 3 | 0 | 0 | 0 |
| Round Total | 9 | 9 | 0 | 0 | 0 |

[^] Results which are Not Assessed should be reviewed by comparing them with the assigned value and other relevant statistics given in the main report. Participants, according to their internal quality criteria, may consider Not Assessed results to be satisfactory, questionable or unsatisfactory. Further information regarding why results may not be assessed is given in the Scheme Information section of the main report.

Please note surplus PT samples are available as QC materials once the round has closed. These samples can be purchased at a reduced rate if you have taken this sample during the main round.

No unsatisfactory results in this round

No questionable results in this round

735 - Meat Indicator Combination

| Analyte | Analyst | Method | Result | llinite | z score (** z' score) | Assigned Value | Ux AV | SDPA | IEXD.SDPA | No of results | Median | Mean | Robust SD | SD |
|--------------------------------|------------|-----------|--------|---------|--------------------------|-------------------|-------|------|-----------|------------------|--------|------|-----------|------|
| Total aerobic mesophilic count | Lab Result | Petrifilm | 79,000 | cfu/g | 0.44 | 55,498 | 0.02 | 0.35 | N/A | 43 | 4.72 | 4.67 | 0.16 | 0.36 |
| Enterobacteriaceae | Lab Result | Petrifilm | 71,800 | cfu/g | 0.56 | 45,749 | 0.03 | 0.35 | N/A | 28 | 4.62 | 4.50 | 0.17 | 0.41 |
| Coliforms | Lab Result | Petrifilm | 50,500 | cfu/g | 0.20 | 43,000 | 0.02 | 0.35 | N/A | 40 | 4.61 | 4.45 | 0.14 | 0.43 |
| Escherichia coli | Lab Result | Petrifilm | 13,000 | cfu/g | 0.03 | 12,703 | 0.02 | 0.35 | N/A | 42 | 4.05 | 3.96 | 0.23 | 0.45 |

736 - Meat Salmonella (P/A)

| Analyte | Test | Analyst | Method | Result | Assigned Value | No of results | Satisfactory % |
|--------------------|--------|------------|--------|----------|----------------|---------------|----------------|
| Salmonella species | Result | Lab Result | Other | Detected | Detected | 65 | 96.9 |

738 - Meat Clost/Staph

| Analyte | Analyst | Method | Result | Units | z score (** z' score) | Assigned Value | Ux AV | SDPA | IEYN SIDPA | No of results | Median | Mean | Robust SD | SD |
|-------------------------------------|------------|----------|--------|-------|--------------------------|-------------------|-------|------|------------|------------------|--------|------|-----------|-----|
| Coagulase positive staphylococci | Lab Result | BP SP 37 | <10 | cfu/g | | Absent | N/A | N/A | N/A | 11 | N/A | N/A | N/A | N/A |

746 - Pseudomonas/Lactic Acid/Yeast/Mould

| Analyte | Analyst | Method | Result | Units | z score (** z' score) | Assigned Value | Ux AV | SDPA | IEYN SDPA | No of results | Median | Mean | Robust SD | SD |
|----------------------|------------|-----------|--------|-------|--------------------------|-------------------|-------|------|-----------|------------------|--------|------|-----------|------|
| Lactic acid bacteria | Lab Result | MRS SP 30 | 72,300 | cfu/g | 0.46 | 50,000 | 0.03 | 0.35 | N/A | 21 | 4.79 | 4.71 | 0.10 | 0.34 |
| Yeast | Lab Result | Petrifilm | 24,000 | cfu/g | 0.58 | 15,000 | 0.04 | 0.35 | N/A | 15 | 4.08 | 4.07 | 0.30 | 0.27 |
| Mould | Lab Result | Petrifilm | <10 | cfu/g | | Absent | N/A | N/A | N/A | 17 | N/A | N/A | N/A | N/A |

** Please note, participant performance for this analyte has been assessed using a z' score, rather than a z score, in order to account for the measurement uncertainty of the assigned value which is not negligible when compared to the SDPA.