External Quality Assessment Scheme

# Gram stain, blood culture Round 1, 2023

#### **Specimens**

The samples S001 and S002 are taken from blood culture bottles and air-fixed on slides.

#### Caution

The slides are unfixed and unstained and should therefore be handled as infectious materials. Fixing kills the microbes.

#### **Examinations**

Gram stain, blood culture

#### **Background information**

Sample S001 (without carbon)
An 82-year-old man. Fever. Peritonitis?

Sample S002 (without carbon)
A 73-year-old woman. Fever. Cholangitis?

# Storage and use

The slides are unfixed and unstained. After arrival, the slides should be stored at +2...8 °C. The slides should be fixed, stained and examined by microscope. The microbes of the specimens are common in clinical specimens, even though they do not grow on all kinds of media.

#### Result reporting

Please enter the results and methods via LabScala (www.labscala.com). Gram staining and Finding parts will be scored. In case of yeast only Finding part will be scored.

S001



S002



#### 2023-01-24

#### **INSTRUCTIONS**

Product no. 5041 LQ760623011-012/FI UN3373

Subcontracting: Sample preparation, sample pretesting

If the kit is incomplete or contains damaged specimens, please report immediately to info@labquality.fi.

The results should be reported no later than **February 21, 2023**.

The expected results of the round are published in LabScala in the View Reports section by February 23, 2023.

#### Inquiries

EQA Coordinator Jaana Paakkanen jaana.paakkanen@labquality.fi

### **Labquality Oy**

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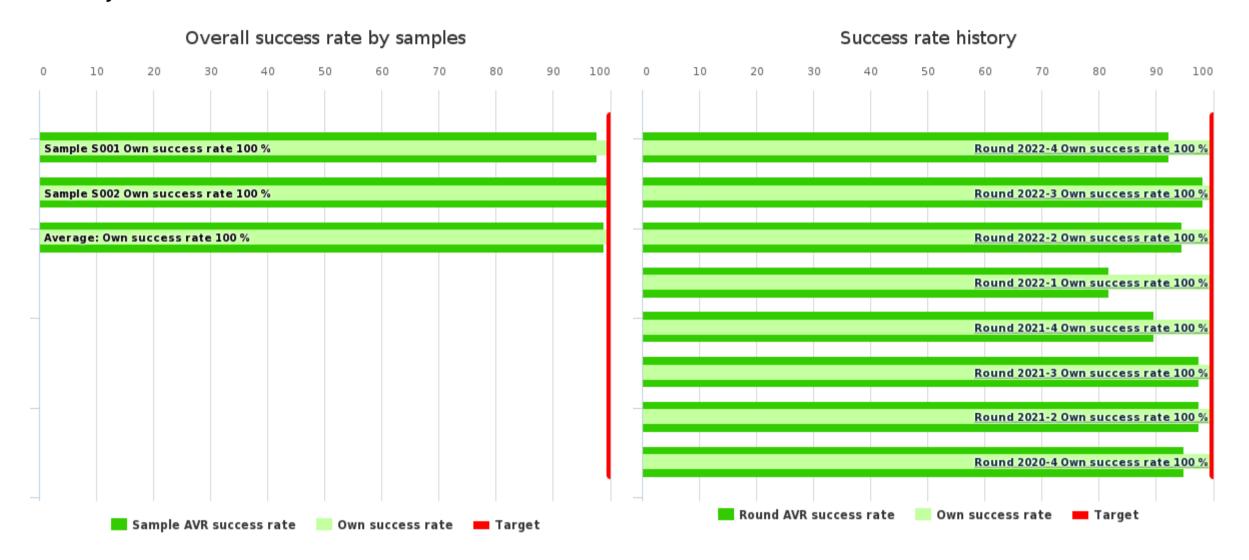




# **Client report**

|  | No of participants | No of responded participants | Response percentage |
|--|--------------------|------------------------------|---------------------|
| Gram stain, blood culture, January, 1-2023 | 108                | 102                          | 94.4 %              |

# **Summary**



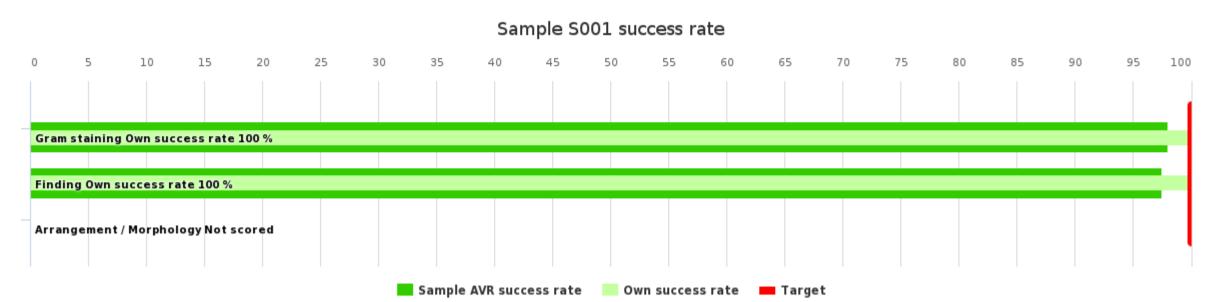
| Summary     | Own score | Max score | Own success rate | Difference | AVR success rate |
|-------------|-----------|-----------|------------------|------------|------------------|
| Sample S001 | 6         | 6         | 100 %            | 2.3 %      | 97.7 %           |
| Sample S002 | 4         | 4         | 100 %            | 0 %        | 100 %            |
| Average:    |           |           | 100 %            | 1.1 %      | 98.9 %           |

| History      | Test nr. | Own success rate | Difference | AVR success rate |
|--------------|----------|------------------|------------|------------------|
| Round 2022-4 | 1        | 100 %            | 7.8 %      | 92.2 %           |
| Round 2022-3 | 1        | 100 %            | 1.7 %      | 98.3 %           |
| Round 2022-2 | 1        | 100 %            | 5.4 %      | 94.6 %           |
| Round 2022-1 | 1        | 100 %            | 18.2 %     | 81.8 %           |
| Round 2021-4 | 1        | 100 %            | 10.4 %     | 89.6 %           |
| Round 2021-3 | 1        | 100 %            | 2.5 %      | 97.5 %           |
| Round 2021-2 | 1        | 100 %            | 2.5 %      | 97.5 %           |
| Round 2020-4 | 1        | 100 %            | 5.1 %      | 94.9 %           |



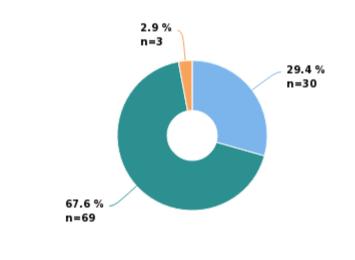


# Sample S001 | Candida albicans, Staphylococcus aureus

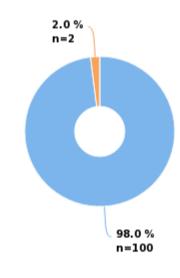


| Sample S001 results | Responded                | Own score | Max score | Own success rate | Difference | AVR success rate | Count |
|---------------------|--------------------------|-----------|-----------|------------------|------------|------------------|-------|
|                     | Gram staining            | 2         | 2         | 100 %            | 2 %        | 98 %             | 204   |
|                     | Finding                  | 4         | 4         | 100 %            | 2.5 %      | 97.5 %           | 204   |
|                     | Arrangement / Morphology | -         | -         | -                | -          | -                | 174   |
| Total               | :                        | 6         | 6         | 100 %            | 2.3 %      | 97.7 %           | 582   |

Sample S001 Gram staining, Candida albicans



Sample S001 Gram staining, Staphylococcus aureus

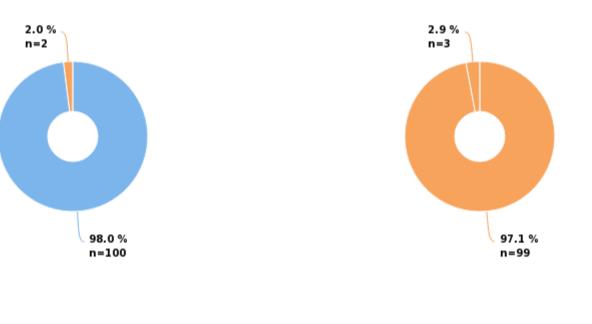


Grampositive No reported finding

Sample S001 Finding, Candida albicans

Yeast No reported finding

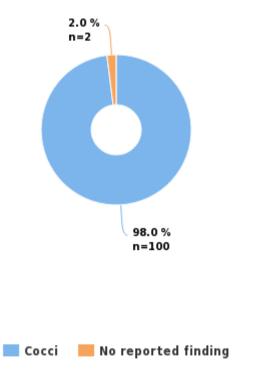
2/6



Sample S001 Finding, Staphylococcus aureus

Grampositive Yeast: no staining result

No reported finding



# **GRAM STAINING**



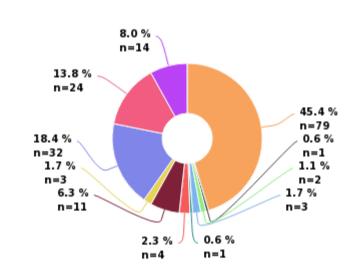


| Finding group         | Result                         | Result<br>count | Own<br>score | Max<br>score | Own<br>success<br>rate | Difference | AVR<br>success<br>rate | Result<br>Score |
|-----------------------|--------------------------------|-----------------|--------------|--------------|------------------------|------------|------------------------|-----------------|
| Candida albicans      |                                | 102             | -            | -            |                        |            | -                      |                 |
|                       | <ul><li>Grampositive</li></ul> | 30              | -            |              |                        |            | -                      | -               |
|                       | Yeast: no staining result      | 69              | -            |              |                        |            | -                      | -               |
|                       | No reported finding            | 3               | -            |              |                        |            | -                      | -               |
| Staphylococcus aureus |                                | 102             | 2            | 2            | 100 %                  | 2 %        | 98 %                   |                 |
|                       | <ul><li>Grampositive</li></ul> | 100             | 2            | 2            | 100 %                  | 0 %        | 100 %                  | 2               |
|                       | No reported finding            | 2               | -            |              |                        |            | 0 %                    | 0               |
| Total:                |                                | 204             | 2            | 2            | 100 %                  | 2 %        | 98 %                   |                 |

# **FINDING**

| Finding group         | Result                  | Result<br>count | Own<br>score | Max<br>score | Own<br>success<br>rate | Difference | AVR<br>success<br>rate | Result<br>Score |
|-----------------------|-------------------------|-----------------|--------------|--------------|------------------------|------------|------------------------|-----------------|
| Candida albicans      |                         | 102             | 2            | 2            | 100 %                  | 2.9 %      | 97.1 %                 |                 |
|                       | <ul><li>Yeast</li></ul> | 99              | 2            | 2            | 100 %                  | 0 %        | 100 %                  | 2               |
|                       | No reported finding     | 3               | -            |              |                        |            | 0 %                    | 0               |
| Staphylococcus aureus |                         | 102             | 2            | 2            | 100 %                  | 2 %        | 98 %                   |                 |
|                       | <ul><li>Cocci</li></ul> | 100             | 2            | 2            | 100 %                  | 0 %        | 100 %                  | 2               |
|                       | No reported finding     | 2               | -            |              |                        |            | 0 %                    | 0               |
| Total:                |                         | 204             | 4            | 4            | 100 %                  | 2.5 %      | 97.5 %                 |                 |

# Sample S001 Arrangement / Morphology





| Arrangement / Morphology | Result                           | Result count |
|--------------------------|----------------------------------|--------------|
|                          | <ul><li>Cocci by group</li></ul> | 79           |
|                          | Cocci in chain                   | 1            |
|                          | Diplococcus                      | 2            |
|                          | <ul><li>Large coccus</li></ul>   | 3            |
|                          | Other interpretation             | 1            |
|                          | Small coccus                     | 4            |
|                          | Tetrads                          | 11           |
|                          | With spores                      | 3            |
|                          | <ul><li>Yeast; Budding</li></ul> | 32           |
|                          | Yeast; Hyphal filaments          | 24           |
|                          | Yeast; Pseudomycelia             | 14           |
|                          | Total:                           | 174          |





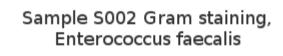
# Sample S002 | Enterococcus faecalis

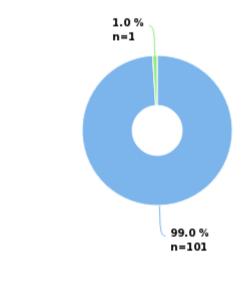


| Sample S002 results | Responded                | Own score | Max score | Own success rate | Difference | AVR success rate | Count |
|---------------------|--------------------------|-----------|-----------|------------------|------------|------------------|-------|
|                     | Gram staining            | 2         | 2         | 100 %            | 0 %        | 100 %            | 104   |
|                     | Finding                  | 2         | 2         | 100 %            | 0 %        | 100 %            | 104   |
|                     | Arrangement / Morphology | -         | -         | -                | -          | -                | 114   |
| Tota                | :                        | 4         | 4         | 100 %            | 0 %        | 100 %            | 322   |

Sample S002 Gram staining, Additional finding

Sample AVR success rate Own success rate Target



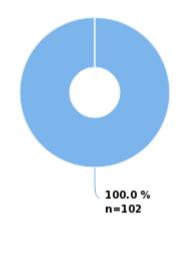






Grampositive



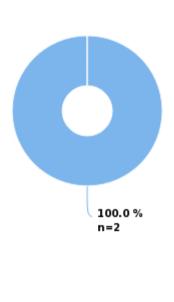


Cocci

4/6



Grampositive Grampositive (weak)



# Cocci

# **GRAM STAINING**



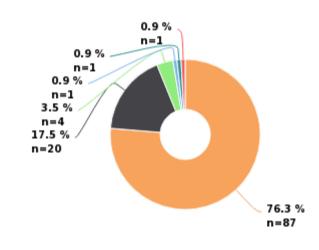


| Finding group         | Result                         | Result<br>count | Own<br>score | Max<br>score | Own<br>success<br>rate | Difference | AVR<br>success<br>rate | Result<br>Score |
|-----------------------|--------------------------------|-----------------|--------------|--------------|------------------------|------------|------------------------|-----------------|
| Enterococcus faecalis |                                | 102             | 2            | 2            | 100 %                  | 0 %        | 100 %                  |                 |
|                       | <ul><li>Grampositive</li></ul> | 101             | 2            | 2            | 100 %                  | 0 %        | 100 %                  | 2               |
|                       | Grampositive (weak)            | 1               | -            |              |                        |            | 100 %                  | 2               |
| Additional finding    |                                | 2               | -            | -            |                        |            | -                      |                 |
|                       | Grampositive                   | 2               | -            |              |                        |            | -                      | -               |
| Total:                |                                | 104             | 2            | 2            | 100 %                  | 0 %        | 100 %                  |                 |

# **FINDING**

| Finding group         | Result                  | Result<br>count | Own<br>score | Max<br>score | Own<br>success<br>rate | Difference | AVR<br>success<br>rate | Result<br>Score |
|-----------------------|-------------------------|-----------------|--------------|--------------|------------------------|------------|------------------------|-----------------|
| Enterococcus faecalis |                         | 102             | 2            | 2            | 100 %                  | 0 %        | 100 %                  |                 |
|                       | <ul><li>Cocci</li></ul> | 102             | 2            | 2            | 100 %                  | 0 %        | 100 %                  | 2               |
| Additional finding    |                         | 2               | -            | _            |                        |            | -                      |                 |
|                       | Cocci                   | 2               | -            |              |                        |            | -                      | -               |
| Total:                |                         | 104             | 2            | 2            | 100 %                  | 0 %        | 100 %                  |                 |

# Sample S002 Arrangement / Morphology





| Arrangement / Morphology | Result                           | Result count |
|--------------------------|----------------------------------|--------------|
|                          | <ul><li>Cocci in chain</li></ul> | 87           |
|                          | Diplococcus                      | 20           |
|                          | Elongated coccus                 | 4            |
|                          | Large coccus                     | 1            |
|                          | Palisade                         | 1            |
|                          | Small coccus                     | 1            |
|                          | Total:                           | 114          |







# **Report Info**

# **PARTICIPANTS**

Altogether 108 laboratories from 11 countries participated in this EQA round.

## **REPORT INFO**

On the front page you can see summaries of overall success rate and sample specific success rates which have been calculated from the scores. The reported results and the scores are presented in the same report but in separate tables.

In general, the expected results are marked with green color. Accepted results may also be indicated with yellow color. Laboratory's own results are indicated with a black radio . In the participant specific report there is also a laboratory specific scoring table for each sample, where you can find your own result and the scores given.

If you have not reported results you will get a note: "You have not responded in time, only global report is available."
For information on report interpretation and performance evaluation, please see the "EQAS Interpretation guidelines" in LabScala User instructions. In case you have any questions regarding the reports, please contact the EQA Coordinator.

# **SCORING**

Results reported for the Gram staining and Finding parts are scored. In case of yeast only Finding part is scored. Scoring is implemented when at least 60% of the participants have reported the correct/expected result and when there are at least three reported results. Laboratory's scores have been converted to percentage (own success rate, % from maximum scores) with a target at 100%. Own success rate is compared with the success rate of all results.

The following general rules are applied:

- 2 points is given to results that are correct/accepted regarding the expected result
  1 point can be given to results that are partly correct/insufficient regarding the expected result
  0 point is given to results that are incorrect/false regarding the expected result

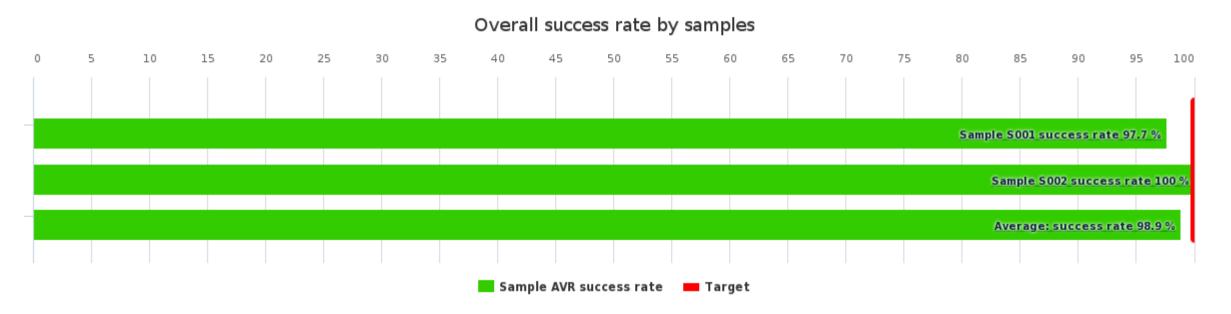
22.03.2023 6/6



# **GLOBAL REPORT**

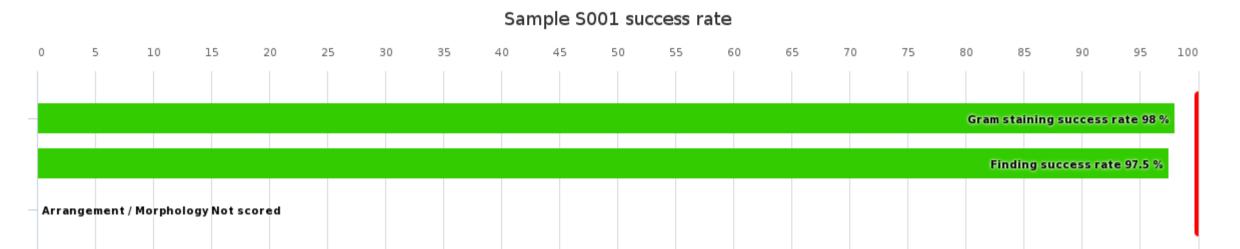
|  | No of participants | No of responded participants | Response percentage |
|--|--------------------|------------------------------|---------------------|
| Gram stain, blood culture, January, 1-2023 | 108                | 102                          | 94.4 %              |

# Summary



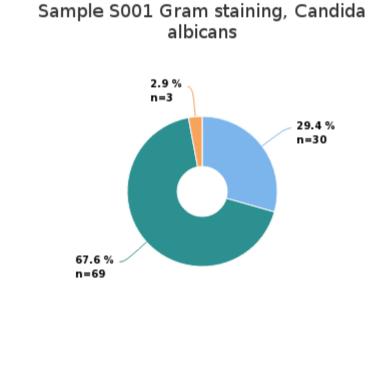
| Summary     | AVR success rate |
|-------------|------------------|
| Sample S001 | 97.7 %           |
| Sample S002 | 100 %            |
| Average:    | 98.9 %           |

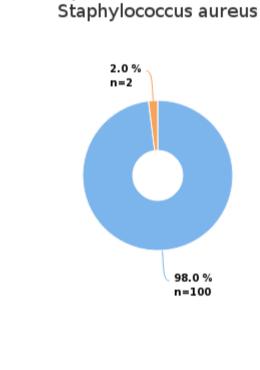
# Sample S001 | Candida albicans, Staphylococcus aureus



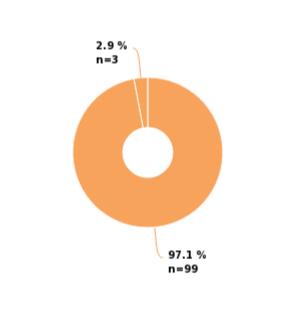
| Sample S001 results | Responded                | AVR success rate | Count |
|---------------------|--------------------------|------------------|-------|
|                     | Gram staining            | 98 %             | 204   |
|                     | Finding                  | 97.5 %           | 204   |
|                     | Arrangement / Morphology | -                | 174   |
|                     | Total:                   | 97.7 %           | 582   |

Sample S001 Gram staining,





Grampositive No reported finding



Yeast No reported finding

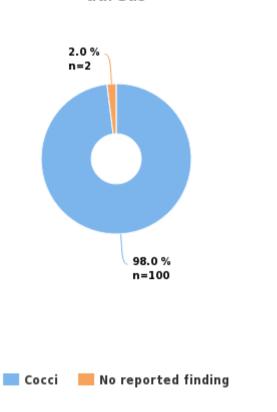
Sample S001 Finding, Candida

albicans

Sample S001 Finding, Staphylococcus aureus

Grampositive Yeast: no staining result

No reported finding



**GRAM STAINING** 

| Tilluling group Result State Result Score | Finding group | Result | Result count | AVR success rate | Result Score |
|---|---------------|--------|--------------|------------------|--------------|
|---|---------------|--------|--------------|------------------|--------------|

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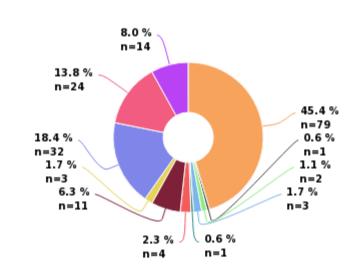
# Gram stain, blood culture, January, 1-2023

| Candida albicans      |                           | 102 | -     |   |
|-----------------------|---------------------------|-----|-------|---|
|                       | Grampositive              | 30  | -     | - |
|                       | Yeast: no staining result | 69  | -     | - |
|                       | No reported finding       | 3   | -     | - |
| Staphylococcus aureus |                           | 102 | 98 %  |   |
|                       | Grampositive              | 100 | 100 % | 2 |
|                       | No reported finding       | 2   | 0 %   | 0 |
| Total:                |                           | 204 | 98 %  |   |

# **FINDING**

| Finding group         | Result              | Result count | AVR success rate | Result Score |
|-----------------------|---------------------|--------------|------------------|--------------|
| Candida albicans      |                     | 102          | 97.1 %           |              |
|                       | Yeast               | 99           | 100 %            | 2            |
|                       | No reported finding | 3            | 0 %              | 0            |
| Staphylococcus aureus |                     | 102          | 98 %             |              |
|                       | Cocci               | 100          | 100 %            | 2            |
|                       | No reported finding | 2            | 0 %              | 0            |
| Total:                |                     | 204          | 97.5 %           |              |

# Sample S001 Arrangement / Morphology

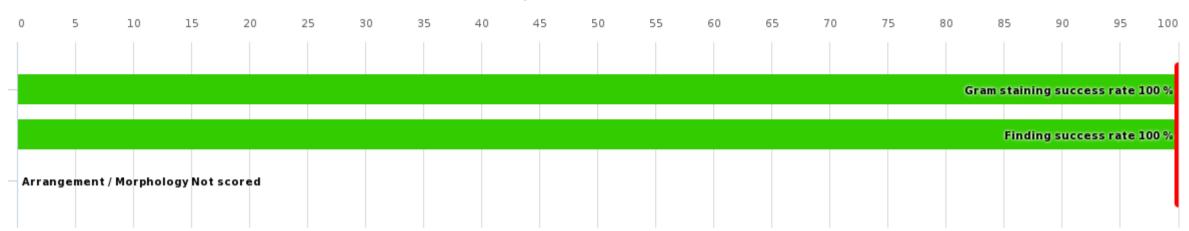




| Arrangement / Morphology | Result                  | Result count |
|--------------------------|-------------------------|--------------|
|                          | Cocci by group          | 79           |
|                          | Cocci in chain          | 1            |
|                          | Diplococcus             | 2            |
|                          | Large coccus            | 3            |
|                          | Other interpretation    | 1            |
|                          | Small coccus            | 4            |
|                          | Tetrads                 | 11           |
|                          | With spores             | 3            |
|                          | Yeast; Budding          | 32           |
|                          | Yeast; Hyphal filaments | 24           |
|                          | Yeast; Pseudomycelia    | 14           |
|                          | Total:                  | 174          |

# Sample S002 | Enterococcus faecalis



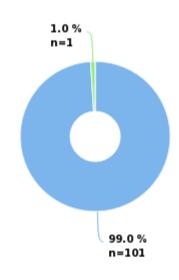


| Sample S002 results | Responded                | AVR success rate | Count |
|---------------------|--------------------------|------------------|-------|
|                     | Gram staining            | 100 %            | 104   |
|                     | Finding                  | 100 %            | 104   |
|                     | Arrangement / Morphology | -                | 114   |
|                     | Total:                   | 100 %            | 322   |

Sample S002 Gram staining,

Additional finding

Sample S002 Gram staining, Enterococcus faecalis

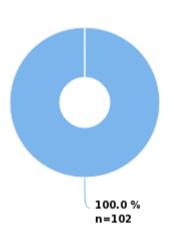






Grampositive



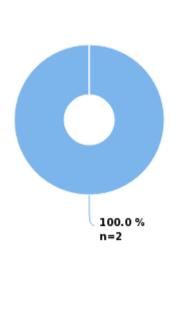


Cocci

4/6



Grampositive Grampositive (weak)



Cocci

# **GRAM STAINING**



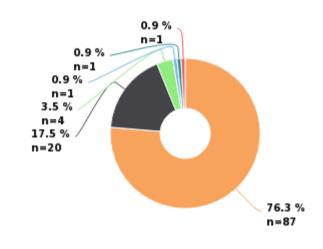
# Gram stain, blood culture, January, 1-2023

| Enterococcus faecalis |                     | 102 | 100 % |   |
|-----------------------|---------------------|-----|-------|---|
|                       | Grampositive        | 101 | 100 % | 2 |
|                       | Grampositive (weak) | 1   | 100 % | 2 |
| Additional finding    |                     | 2   | -     |   |
|                       | Grampositive        | 2   | -     | - |
| Total:                |                     | 104 | 100 % |   |

# **FINDING**

| Finding group         | Result | Result count | AVR success rate | Result Score |
|-----------------------|--------|--------------|------------------|--------------|
| Enterococcus faecalis |        | 102          | 100 %            |              |
|                       | Cocci  | 102          | 100 %            | 2            |
| Additional finding    |        | 2            | -                |              |
|                       | Cocci  | 2            | -                | -            |
| Total:                |        | 104          | 100 %            |              |

# Sample S002 Arrangement / Morphology





| Arrangement / Morphology | Result           | Result count |
|--------------------------|------------------|--------------|
|                          | Cocci in chain   | 87           |
|                          | Diplococcus      | 20           |
|                          | Elongated coccus | 4            |
|                          | Large coccus     | 1            |
|                          | Palisade         | 1            |
|                          | Small coccus     | 1            |
|                          | Total:           | 114          |



# Gram stain, blood culture, January, 1-2023

# **Report Info**

# **PARTICIPANTS**

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## **REPORT INFO**

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21.03.2023 6/6

**External Quality Assessment Scheme** 

# Gram stain, blood culture Round 1, 2023

#### **Specimens**

The samples were three air-fixed microbial suspensions on a slide. The sample lots were tested in an accredited Finnish reference laboratory. Based on the quality controls conducted by the sample material manufacturer, pretesting and the results obtained in the round, the sample lots are to be considered as homogeneous, stable and suitable for external quality assessment. The materials were sent without temperature control packaging.

The content of the samples was as follows:

Sample S001 (LQ760623011)

Candida albicans, yeast

Staphylococcus aureus, grampositive cocci

Background information: An 82-year-old man. Fever. Peritonitis?

Sample S002 (LQ760623012)

Enterococcus faecalis, grampositive cocci

Background information: A 73-year-old woman. Fever. Cholangitis?

## Report info

Please see the description of the data analysis on the last page of the laboratory-specific reports and global reports. It is important to read the Final report first, because it contains important information of the samples and results in each round.

#### **Comments - Expert**

Average success rate in this round was 98.9%.

# Sample S001 Candida albicans and Staphylococcus aureus

Candida albicans yeast cells are spherical or ovoid (2.0–7.0 x 3.0–8.5  $\mu$ m). Yeast cells appear alone or with a budding cell. Many laboratories (41.3%) commented the appearance of the daughter cells. In addition, pseudohyphae form of the cells exist. 97.1% of the participating laboratories reported finding correctly as a yeast.

Along with yeast there was another microbe in the sample. Staphylococcus aureus is a grampositive coccus. The cells are small (1  $\mu$ m in diameter) and spherical. The cells occur in group. 98.0% of the participating laboratories reported grampositive coccus and 79.0% of them commented the arrangement of the cells in groups.

Including both findings, an average success rate for this sample was 97.7%.

#### Sample S002 Enterococcus faecalis

Enterococcus faecalis is a grampositive coccus. The cells are spherical to ovoid  $0.5-1.0~\mu m$  in diameter. The cells arrange normally in pairs or in short chains. All participating laboratories reported the gramstaining correctly as grampositive coccus. 76.3% of the laboratories commented the arrangement of the cells in chains.

### **Exceptions in scoring**

No exceptions.

#### 2023-03-21

#### FINAL REPORT

Product no. 5041

Subcontracting: Sample preparation, Sample pretesting

 Samples sent
 2023-01-24

 Round closed
 2023-02-21

 Expected results
 2023-02-23

 Final report
 2023-03-21

#### Request for correction

Typing errors in laboratory's result forms are on laboratory's responsibility. Labquality accepts responsibility only for result processing. Requests must be notified by writing within three weeks from the date of this letter.

#### Authorized by

EQA Coordinator Jaana Paakkanen jaana.paakkanen@labquality.fi

#### **Expert**

PhD, Clinical Microbiologist, Päivi Mähönen, Vita Laboratories, Helsinki, Finland

#### **Labquality Oy**

Kumpulantie 15 FI-00520 HELSINKI Finland

Tel. + 358 9 8566 8200 Fax + 358 9 8566 8280

info@labquality.fi www.labquality.com



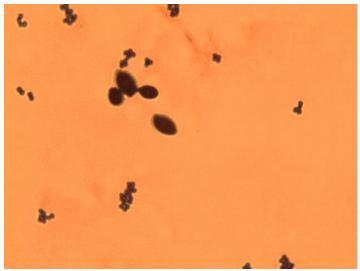


# **End of report**

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# 5041 Gram stain, blood culture 1, 2023 5041 Gramvärjäys, veriviljely 1, 2023



# S001 Candida albicans

#### a yeast

Characteristics: small, ovoid yeast cells alone and with a daughter cell, pseudohyphae.

### hiiva

Tunnusomaista: pieni, soikea hiivasolu, yksin ja yhdessä tytärsolujen kanssa, valerihmaa.



# S001 Staphylococcus aureus

## a grampositive cocci

Characteristics: small coccus in groups.

## grampositiivinen kokki

Tunnusomaista: pieniä ryhmäkokkeja.



# S002 Enterococcus faecalis

## a grampositive cocci

Characteristics: spherical or ovoid coccus in short chains and in pairs.

# grampositiivinen kokki

Tunnusomaista: pyöreä tai soikea kokki, esiintyy lyhyissä ketjuissa ja pareittain.



