

Aerobiology Laboratory - VA
43760 Trade Center Place
Sterling, Virginia 20166
Attn: Chris Blevins
Project: **MRSA Test**
Condition of Sample(s) Upon Receipt: Acceptable

Date Collected: 03/16/2021
Date Received: 03/17/2021
Date Analyzed: 03/19/2021
Date Reported: 03/20/2021
Project ID: 21021137
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Client Sample #: 1
Sample Location: OR Bench
Test: 1121, Air, MRSA and Total Bacterial Count w/ID's
Positive Hole Corrected Result: **26 CFU/m³**

Lab Sample #: 21021137-001
Positive Hole: **219**
Air Volume: **200 (L)**
MRL: **5**

Organism(s) Isolated:	Raw Count	CFU/m ³	% Total	Reservoirs
Methicillin-resistant Staphylococcus aureus-MRSA	2	10	40	Human
Micrococcus species	3	15	60	Human
	5	25	~100%	

Client Sample #: 2
Sample Location: OR Room 4
Test: 1121, Air, MRSA and Total Bacterial Count w/ID's
Positive Hole Corrected Result:

Lab Sample #: 21021137-002
Positive Hole: **219**
Air Volume: **200 (L)**
MRL: **5**

Comments: No MRSA isolated.

Client Sample #: 3
Sample Location: OR Bench
Test: 1122, Surface / Wipe MRSA and Total Bacterial Count w/ID's
Results: **50 CFU/in²**

Lab Sample #: 21021137-003
Area: **1 (in²)**

Organism(s) Isolated:	Raw Count	CFU/in ²	% Total	MRL
Methicillin-resistant Staphylococcus aureus-MRSA	4	40	80	10
Micrococcus species	1	10	20	10
	5	50	~100%	

Client Sample #: 4
Sample Location: OR Room 4
Test: 1122, Surface / Wipe MRSA and Total Bacterial Count w/ID's
Results:

Lab Sample #: 21021137-004
Area: **1 (in²)**
MRL: **10**

Comments: No MRSA isolated.

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Client Sample #: 5
 Sample Location: OR 1
 Test: 1123, Bulk / Dust MRSA and Total Bacterial Count w/ID's
 Results: **40 CFU/g**

Lab Sample #: 21021137-005

Mass Analyzed: **0.5 (g)**

Organism Name:	Raw Count	CFU/g	% Total	Reservoirs
Coag-negative Staphylococcus species	2	20	50	Human
Methicillin-resistant Staphylococcus aureus-MRSA	2	20	50	Human
	4	40	~100%	

Client Sample #: 6
 Sample Location: OR 4
 Test: 1123, Bulk / Dust MRSA and Total Bacterial Count w/ID's
 Results:

Lab Sample #: 21021137-006

Mass Analyzed: **0.5 (g)**

Comments: No MRSA isolated.

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Footnotes and Additional Report Information

Debris Rating Table

1	Minimal (<5%) particulate present	Reported values are minimally affected by particulate load.
2	5% to 25% of the trace occluded with particulate	Negative bias is expected. The degree of bias increases directly with the percent of the trace that is occluded.
3	26% to 75% of the trace occluded with particulate	Negative bias is expected. The degree of bias increases directly with the percent of the trace that is occluded.
4	75% to 90% of the trace occluded with particulate	Negative bias is expected. The degree of bias increases directly with the percent of the trace that is occluded.
5	Greater than 90% of the trace occluded with particulate	Quantification not possible due to large negative bias. A new sample should be collected at a shorter time interval or other measures taken to reduce particulate load.

Aerobiology Laboratory Associates, Inc. shall be responsible for all the information provided in the report, except when information is provided by the customer. Data provided by a customer can affect the validity of results and shall be clearly identified. Results apply to the samples as received. Aerobiology Laboratory Associates, Inc. is not responsible for the sampling activity, such as air and water volume, area, and mass unit. The report shall not be reproduced except in full without the approval of the laboratory to ensure that parts of a report are not taken out of context. Data interpretation of this report will be the client responsibility based on their sampling.

1. Penicillium/Aspergillus group spores are characterized by their small size, round to ovoid shape, being unicellular, and usually colorless to lightly pigmented. There are numerous genera of fungi whose spore morphology is similar to that of the Penicillium/Aspergillus type. Two common examples would be Paecilomyces and Acremonium. Although the majority of spores placed in this group are Penicillium, Aspergillus, or a combination of both. Keep in mind that these are not the only two possibilities.

2. Ascospores are sexually produced fungal spores formed within an ascus. An ascus is a sac-like structure designed to discharge the ascospores into the environment, e.g. Ascobolus.

3. Basidiospores are typically blown indoors from outdoors and rarely have an indoor source. However, in certain situations a high basidiospore count indoors may be indicative of a wood decay problem or wet soil.

4. The colorless group contains colorless spores which were unidentifiable to a specific genus. Examples of this group include Acremonium, Aphanocladium, Beauveria, Chrysosporium, Engyodontium microconidia, yeast, some arthrospores, as well as many others.

5. Hyphae are the vegetative mode of fungi. Hyphal elements are fragments of individual Hyphae. They can break apart and become airborne much like spores and are potentially allergenic. A mass of hyphal elements is termed the mycelium. Hyphae in high concentration may be indicative of colonization.

6. Dash (-) in this report, under raw count column means 'not detected (ND)'; otherwise 'not applicable' (NA).

7. The positive-hole correction factor is a statistical tool which calculates a probable count from the raw count, taking into consideration that multiple particles can impact on the same hole; for this reason the sum of the calculated counts may be less than the positive hole corrected total.

8. Due to rounding totals may not equal 100%.

9. Analytical Sensitivity for each spores is different for Non-viable sample when the spores are read at different percentage. Analytical Sensitivity is calculated as spr/m^3 divided by raw count. $spr/m^3 = \text{raw counts} \times (100/\% \text{ read}) \times (1000/\text{Sample volume})$. If Analytical Sensitivity is 13 spr/m^3 at 100% read, Analytical Sensitivity at 50% read would be 27 spr/m^3 , which is 2 times higher. Analytical Sensitivity provided on the report is based on an assumed 100% of the trace being analyzed.

10. Minimum Reporting Limits (MRL) for BULKs, DUSTS, SWABS, and WATER samples are a calculation based on the sample size and the dilution plate on which the organism was counted. Results are a compilation of counts taken from multiple dilutions and multiple medias. This means that every genus of fungi or bacteria recovered can be counted on the plate on which it is best represented.

11. If the final quantitative result is corrected for contamination based on the blank, the blank correction is stated in the sample comments section of the report.

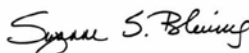
12. The results in this report are related to this project and these samples only.

13. For samples with an air volume of < 100L, the number of significant figures in the result should be considered (2) two. For samples with air volumes between 100-999L, the number of significant figures in the result should be considered (3) three. For example, a sample with a result of 55,443 spr/m^3 from a 75L sample using significant figures should be considered 55,000. The same result of 55,443 from a 150L sample using significant figures should be considered 55,400 spr/m^3 .

14. If the In/Out ratio is greater than 100 times it is indicated >100/1, rather than showing the real value.

Terminology Used in Direct Exam Reporting

Conidiophores are a type of modified hyphae from which spores are born. When seen on a surface sample in moderate to numerous concentrations they may be indicative of fungal growth.



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Laboratory Director