

Report for:

Mr. Randy Mount Ram Restoration 2800 East River Road Suite A Dayton, OH 45439

Regarding: Project: Waynesville Highschool; 4/21 Post test 2

EML ID: 3010549

Approved by:

Technical Manager Francina Thadigiri Dates of Analysis:

MoldReport Culturable air fungi (Incl. Asp spp.): 08-31-2022

Service SOPs: MoldReport Culturable air fungi (Incl. Asp spp.) (EM-MY-S-1043) AIHA-LAP, LLC accredited service, Lab ID #179623

All samples were received in acceptable condition unless noted in the Report Comments portion in the body of the report. Due to the nature of the analyses performed, field blank correction of results is not applied. The results relate only to the samples as received and tested. Information supplied by the client which can affect the validity of results: sample air volume.

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Eurofins EMLab P&K

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Client: Ram Restoration

C/O: Mr. Randy Mount

Re: Waynesville Highschool; 4/21 Post test 2

Date of Sampling: 04-21-2022

Date of Receipt: 08-23-2022

Date of Report: 08-31-2022

CULTURABLE AIR FUNGI REPORT

| Location: | 01: computer lab 2 | | .02: | | 03: | | 04: | |
|----------------------------|-----------------------|---------|--------------|---------|-----------------------|---------|------------|---------|
| | | | media center | | first floor left hall | | Room 16 | |
| Comments (see below) | None | | None | | None | | None | |
| Lab ID-Version‡: | 14494600-1 | | 14494601-1 | | 14494602-1 | | 14494603-1 | |
| Analysis Date: | 08/31/2022 | | 08/31/2022 | | 08/31/2022 | | 08/31/2022 | |
| Medium: | MEA | | MEA | | MEA | | MEA | |
| | raw ct. | cfu*/m3 | raw ct. | cfu*/m3 | raw ct. | cfu*/m3 | raw ct. | cfu*/m3 |
| Aspergillus niger | | | | | | | | |
| Aspergillus ochraceus | | | | | | | | |
| Aspergillus versicolor | | | | | | | | |
| Aureobasidium | | | | | | | | |
| Basidiomycetes | | | | | | | | |
| Bipolaris/Drechslera group | | | | | | | | |
| Botrytis | | | | | | | | |
| Chaetomium | | | | | | | | |
| Cladosporium | | | | | | | | |
| Curvularia | | | | | | | | |
| Epicoccum | | | | | | | | |
| Fusarium | | | | | | | | |
| Non-sporulating fungi | | | | | | | | |
| Paecilomyces | | | | | | | | |
| Penicillium | | | | | | | | |
| Phoma | | | | | | | | |
| Rhizopus | | | | | | | | |
| Stachybotrys chartarum | | | | | | | | |
| Ulocladium | | | | | | | | |
| Yeasts | | | | | | | | |
| Positive Hole | 0 | | 0 | | 0 | | 0 | |
| Sample volume (liters) | 105 | | 105 | | 105 | | 105 | |
| § TOTAL CFU*/M3 | | < 10 | | < 10 | | < 10 | | < 10 |

^{*} cfu = colony forming units

Positive hole correction chart used for all calculations

Comments:

Note: Interpretation is left to the company and/or persons who conducted the field work. Variation is an inherent part of biological sampling. The presence or absence of a few genera in small numbers should not be considered abnormal.

NORMAL SPORE LEVELS: Indoor spore levels usually average 30 to 80% of the outdoor spore level at the time of sampling, with the same general distribution of spore types. Filtered air, air-conditioned air, or air remote from outside sources may average 5 to 15% of the outside air at the time of sampling. (These percentages are guidelines, only. A major factor is the accessibility of outdoor air. A residence with open doors and windows and heavy foot traffic may average 95% of the outdoor level while high rise office buildings with little air exchange may average 2%. Dusty interiors may exceed 100% of the outdoors to some degree, but will still mirror the outdoor distribution of spore types.)

PROBLEM INTERIORS: A substantial increase of one or two spore types which are inconsistent with and non-reflective of the outside distribution of spore types is usually indicative of an indoor reservoir of mold growth.

The limit of detection is 1 raw count per volume of air sampled. The analytical sensitivity is 1 raw count/volume x the positive hole correction factor.

Fungal culture types listed without a count or data entry were not detected during the course of the analysis for the respective sample.

 $[\]ddagger$ A "Version" indicated by -"x" after the Lab ID# with a value greater than 1 indicates a sample with amended data. The revision number is reflected by the value of "x".

[§] Total CFU/m3 has been rounded to two significant figures to reflect analytical precision.