



**Customer Name:** AGX, Inc. **Sample Date:** August 19, 2019  
**Customer Address:** 207 Pine Creek Road **Date Received:** August 20, 2019  
 Wexford, PA 15090 **Date of Report:** August 21, 2019

**Customer Phone:** (724) 934-4249 **Fax:** (724) 934-5677  
**PO Number:** **Attention:** Amber Brancolini  
**Project Name/Number:** Summit Township Elementary

Customer sample numbers below are uniquely identified by prefixing Laboratory # 88218-19

Airborne Spore Trap Analysis - AllergencoD												
Analytical Method: USMS-M008												
Total Volume (L)	75				75				75			
Sample Number	STE-10-01				STE-2-02				STE-4-03			
Location:	Room 10				Room 2				Room 4			
Particle ID	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%
Alternaria												
Ascospores	17	13	221	1%	14	13	182	11%	14	13	182	0%
Aspergillus/Penicillium-like	112	178	19,936	95%	31	13	403	25%	139	267	37,113	94%
Basidiospores	5	13	65	0%	22	13	286	18%	9	13	117	0%
Bipolaris/Drechslera												
Cercospora												
Chaetomium												
Cladosporium	64	13	832	4%	55	13	715	45%	170	13	2,210	6%
Curvularia												
Epicoccum												
Helicomyces												
Nigrospora												
Oidium												
Pithomyces/Ulocladium	2	13	26	0%					1	13	13	0%
Polythrincium												
Rusts												
Smuts/ Myxomycetes												
Stachybotrys												
Torula												
Trichoderma												
Unidentified dematiaceous conidia												
Unidentified hyaline conidia												
<b>Total Mold (Spores/m<sup>3</sup> of air)</b>	200		21,080		122		1,586		333		39,635	
Pollen	0	13	< 13		0	13	< 13		0	13	< 13	
Hyphal Fragments	4	13	52		2	13	26		2	13	26	
Insect Fragments												
Plant Fragments												
Skin Cell Fragments			1				1				1	
Debris			2				1				1	
Analyst Initials			HC				HC				HC	
Date Analyzed			08/20/19				08/20/19				08/20/19	
Cassette Serial # / Exp Date:			2796548 09/2019				2796542 09/2019				2796562 09/2019	

Entire trace analyzed. Results relate only to the samples tested. Results are reported as calculated. For biological data, the first and/or second digit should be considered significant. Total percentage may not equal 100% due to rounding. Percentages reported as 0% are greater than 0 and less than 0.5%. The *Aspergillus/Penicillium*-like category cannot be differentiated by non-viable sampling methods.  
 AS=Analytical Sensitivity (spore/m<sup>3</sup>); Blank Lines = None Detected

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Technical Manager:

*Sharon Fanchalsky*

Sharon Fanchalsky, AS, MLT (ASCP)



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Airborne Spore Trap Analysis - AllergencoD												
Analytical Method: USMS-M008												
Total Volume (L)	75				75				75			
Sample Number	STE-1-04				STE-5-05				STE-6-06			
Location:	Room 1				Room 5				Room 6			
Particle ID	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%
Alternaria												
Ascospores	3	13	39	4%	4	13	52	1%	7	13	91	6%
Aspergillus/Penicillium-like	38	13	494	55%	367	13	4,771	94%	75	13	975	62%
Basidiospores	14	13	182	20%	2	13	26	1%	2	13	26	2%
Bipolaris/Drechslera												
Cercospora												
Chaetomium												
Cladosporium	14	13	182	20%	19	13	247	5%	37	13	481	31%
Curvularia												
Epicoccum												
Helicomyces												
Nigrospora												
Oidium												
Pithomyces/Ulocladium												
Polythrincium												
Rusts												
Smuts/ Myxomycetes												
Stachybotrys												
Torula												
Trichoderma												
Unidentified dematiaceous conidia												
Unidentified hyaline conidia												
<b>Total Mold (Spores/m<sup>3</sup> of air)</b>	69		897		392		5,096		121		1,573	
Pollen	0	13	< 13		0	13	< 13		0	13	< 13	
Hyphal Fragments												
Insect Fragments												
Plant Fragments												
Skin Cell Fragments			1				1				1	
Debris			2				2				2	
Analyst Initials	LS				LS				LS			
Date Analyzed	08/20/19				08/20/19				08/20/19			
Cassette Serial # / Exp Date:	2796547 09/2019				2796552 09/2019				2796553 09/2019			

Entire trace analyzed. Results relate only to the samples tested. Results are reported as calculated. For biological data, the first and/or second digit should be considered significant. Total percentage may not equal 100% due to rounding. Percentages reported as 0% are greater than 0 and less than 0.5%. The *Aspergillus/Penicillium*-like category cannot be differentiated by non-viable sampling methods.  
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Airborne Spore Trap Analysis - AllergencoD												
Analytical Method: USMS-M008												
Total Volume (L)	75				75				75			
Sample Number	STE-7-07				STE-13-08				STE-16-09			
Location:	Room 7				Room 13				Room 16			
Particle ID	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%
Alternaria												
Ascospores	26	13	338	4%	18	13	234	6%	26	13	338	12%
Aspergillus/Penicillium-like	480	13	6,240	83%	165	13	2,145	58%	111	13	1,443	50%
Basidiospores	29	13	377	5%	28	13	364	10%	36	13	468	16%
Bipolaris/Drechslera												
Cercospora												
Chaetomium												
Cladosporium	46	13	598	8%	71	13	923	25%	51	13	663	23%
Curvularia												
Epicoccum					1	13	13	0%				
Helicomyces					1	13	13	0%				
Nigrospora												
Oidium												
Pithomyces/Ulocladium												
Polythrincium												
Rusts												
Smuts/ Myxomycetes					2	13	26	1%				
Stachybotrys												
Torula												
Trichoderma												
Unidentified dematiaceous conidia												
Unidentified hyaline conidia												
<b>Total Mold (Spores/m<sup>3</sup> of air)</b>	<b>581</b>		<b>7,553</b>		<b>286</b>		<b>3,718</b>		<b>224</b>		<b>2,912</b>	
Pollen	1	13	13		0	13	< 13		0	13	< 13	
Hyphal Fragments												
Insect Fragments	1	13	13		6	13	78		1	13	13	
Plant Fragments												
Skin Cell Fragments			1				1				1	
Debris			1				1				1	
Analyst Initials	BM				BM				BM			
Date Analyzed	08/20/19				08/20/19				08/20/19			
Cassette Serial # / Exp Date:	2796558 09/2019				2796567 09/2019				2796569 09/2019			

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Airborne Spore Trap Analysis				- AllergencoD			
Analytical Method:				USMS-M008			

Total Volume (L)	75				75				75			
	STE-17-10				STE-Gd-11				STE-Prm-12			
Location:	Room 17				Guidance				Principals Office			
Particle ID	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%
Alternaria												
Ascospores	13	13	169	13%	2	13	26	0%				
Aspergillus/Penicillium-like	46	13	598	44%	145	178	25,810	99%	40	13	520	91%
Basidiospores	39	13	507	38%					3	13	39	7%
Bipolaris/Drechslera												
Cercospora												
Chaetomium												
Cladosporium	6	13	78	6%	14	13	182	1%	1	13	13	2%
Curvularia												
Epicoccum												
Helicomyces												
Nigrospora												
Oidium												
Pithomyces/Ulocladium												
Polythrincium												
Rusts												
Smuts/ Myxomycetes												
Stachybotrys												
Torula												
Trichoderma												
Unidentified dematiaceous conidia												
Unidentified hyaline conidia												
<b>Total Mold (Spores/m<sup>3</sup> of air)</b>	104		1,352		161		26,018		44		572	
Pollen	0	13	< 13		0	13	< 13		0	13	< 13	
Hyphal Fragments												
Insect Fragments												
Plant Fragments												
Skin Cell Fragments			1				1				1	
Debris			2				2				2	
Analyst Initials			LS				LS				LS	
Date Analyzed			08/20/19				08/20/19				08/20/19	
Cassette Serial # / Exp Date:			2796581 09/2019				2796574 09/2019				2796576 09/2019	

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**Airborne Spore Trap Analysis - AllergencoD**  
**Analytical Method: USMS-M008**

Total Volume (L)		75											
Sample Number		STE-Out-13											
Location:		Outside											
Particle ID	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	Raw ct.	AS	Spores/m <sup>3</sup>	%	
Alternaria	1	13	13	0%									
Ascospores	128	13	1,664	26%									
Aspergillus/Penicillium-like	24	13	312	5%									
Basidiospores	172	13	2,236	36%									
Bipolaris/Drechslera													
Cercospora													
Chaetomium													
Cladosporium	145	13	1,885	30%									
Curvularia													
Epicoccum	1	13	13	0%									
Helicomyces	2	13	26	0%									
Nigrospora													
Oidium													
Pithomyces/Ulocladium	3	13	39	1%									
Polythrincium													
Rusts	2	13	26	0%									
Smuts/ Myxomycetes	6	13	78	1%									
Stachybotrys													
Torula													
Trichoderma													
Unidentified dematiaceous conidia													
Unidentified hyaline conidia													
Total Mold (Spores/m <sup>3</sup> of air)	484		6,292										
Pollen	1	13	13										
Hyphal Fragments	1	13	13										
Insect Fragments													
Plant Fragments													
Skin Cell Fragments			1										
Debris			1										
Analyst Initials			HC										
Date Analyzed			08/20/19										
Cassette Serial # / Exp Date:			2931337 03/2020										

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## SPORE TRAP INTERPRETATION TIPS

Currently there are no numeric standards for indoor airborne or surface microbial contamination. Suggested guidelines are constantly being reviewed and updated as more information is collected.

Some common denominators should be considered when interpreting results:

1. Comparison of indoor/outdoor concentration ratios.
2. Complaint vs. non-complaint areas or affected vs. non-affected areas.
3. Consider air exchange rates and activity levels in a building structure, weather, and season of the year.
4. Rank order assessment and concentration (e.g. Spores/m<sup>3</sup> of air) of the fungi.
5. Predominant fungal genera: Are there water indicator microorganisms present, such as but not limited to: *Chaetomium*, *Stachybotrys*, *Rhodotorula*, *Trichoderma*, and *Scopulariopsis*.
6. Generally the fungal counts indoors should be lower than outdoor counts and the types of fungi found indoors should be similar to outdoors.
7. There is always a potential bias from infiltration of outdoor air, poor housekeeping, excessive indoor relative humidity, or potential contamination sources (e.g. water intrusion through a basement wall) that may negatively influence post remedial verification (PRV) or clearance levels.
8. The investigator should look for various patterns among the indoor types of molds detected:
  - a. Increased levels of primary (1st) colonizers in damp or moisture intrusion areas of homes or commercial buildings: ***Aspergillus/Penicillium*** or ***Cladosporium*** are usually noted.
  - b. ***Chaetomium*** or ***Stachybotrys*** are tertiary (3rd) colonizers of indoor materials and are usually associated with chronic long standing water/moisture issues in a building.
  - c. The presence of ***hyphal fragments*** or ***fruiting structures*** noted on spore trap samples usually indicates amplification (growth) of fungi on building substrates.
  - d. ***Ascospores*** and ***basidiospores*** noted on indoors spore trap samples most often represent the entrance of inadequately filtered outdoor air. During inclement weather, remember to note time, temperature, and season. Most indoor materials will not support the growth of these fungi.
9. When unidentified hyaline (clear) or dematiaceous (dark-pigmented) conidia are noted on a spore trap sample, it indicates that no particular fungus can be identified. These fungal conidia may represent such yeast-like fungi as *Aureobasidium*, *Sporidiobolus*, unidentifiable *Acremonium* species, Basidiomycetes (basidiospores), and Ascomycetes (ascospores).
10. Keep in mind when interpreting spore trap sample reports, that indoor levels may be higher than corresponding outdoor levels (winter time in the Northern U.S.) with a predominance of *Aspergillus/Penicillium* or *Cladosporium* conidia with no significant amplification of any molds.

## SPORE TRAP GUIDELINES FOR INDOOR MICROBIAL CONTAMINATION

<b>DEBRIS RATING for SPORE TRAP ANALYSIS (using 600X magnification)</b> (Air-O-Cell, Micro 5, Allergenco D, Cyclex d, VersaTrap, etc.)		
<b>DEBRIS RATING</b>	<b>CONDITIONS FOR REPORTING DEBRIS RATING</b>	<b>SIGNIFICANCE</b>
0	A visible trace, including particulates and debris, is not observed.	Indicates the sample was a blank, the area is exceptionally clean, or improper sampling occurred.
1	Debris is present and <10% of the average viewing field is obscured.	Minimal amount of debris is observed.
2	Debris is present and 10% to <40% of the average viewing field is obscured.	Low amount of debris is observed, counts may be affected.
3*	Debris is present and 40% to 75% of the average viewing field is obscured.	Moderate amount of debris is observed, counts of conidia/hyphal fragments may be underestimated.
4*	Debris is present and >75% of the average viewing field is obscured.	High amount of debris is observed, counts are estimated.
5* See Relative Abundance chart below	Excessive debris is present	Periphery of trace analyzed. Relative amounts of conidia/hyphal fragments noted. Suggest recollection.
6	Slide completely obscured by excessive debris.	Unable to analyze. Recollect sample.

\* A rating of 3 or greater indicates that the accuracy of the analysis is likely affected.

<b>RELATIVE ABUNDANCE of OBSERVED CONIDIA &amp; HYPHAL FRAGMENTS</b>	
<b>RATING</b>	<b>Relative Amounts of Observed Fungal Structures per high power field (600X)</b>
Rare	0-1
Few	2 to 5
Moderate	6 to 10
Many	11 to 100
Numerous	>100

<b>SKIN CELL ANALYSIS</b>	
<b>SKIN CELL RATING</b>	<b>Relative Amounts of Observed Skin Cells per high power field (600X)</b>
0	No skin cells present
1	0-1
2	2 to 5
3	6 to 10
4	11 to 15
5	≥16

**\*End of Report\***