

### EMLab P&K

Report for:

Mr. Brian Poplarchick LaBella Associates, D.P.C 1000 Dunham Drive, Suite B Dunmore, PA 18512

Regarding:

Project: ESASD- 2200223; ESASD- East Stroudsburg H.S. South

EMĹ ID: 2340919

Approved by:

Dates of Analysis:

Spore trap analysis: 01-27-2020

Technical Manager Ariunaa Jalsrai

Service SOPs: Spore trap analysis (EM-MY-S-1038) AIHA-LAP, LLC accredited service, Lab ID #103005

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. Sample air volume is supplied by the client.

Eurofins EMLab P&K ("the Company") shall have no liability to the client or the client's customer with respect to decisions or recommendations made, actions taken or courses of conduct implemented by either the client or the client's customer as a result of or based upon the Test Results. In no event shall the Company be liable to the client with respect to the Test Results except for the Company's own willful misconduct or gross negligence nor shall the Company be liable for incidental or consequential damages or lost profits or revenues to the fullest extent such liability may be disclaimed by law, even if the Company has been advised of the possibility of such damages, lost profits or lost revenues. In no event shall the Company's liability with respect to the Test Results exceed the amount paid to the Company by the client therefor.

Eurofins EMLab P&K's LabServe® reporting system includes automated fail-safes to ensure that all AIHA-LAP, LLC quality requirements are met and notifications are added to reports when any quality steps remain pending.

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Client: LaBella Associates, D.P.C Date of Sampling: 01-23-2020 C/O: Mr. Brian Poplarchick Date of Receipt: 01-24-2020 Re: ESASD- 2200223; ESASD- East Stroudsburg H. Date of Report: 01-28-2020

S. South

### SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	001: E. Stroudsburg High School South- Main Office			E. Stroudsburg High School South- Rm.		
Comments (see below)		None		A		
Lab ID-Version‡:		11146288-	1		11146289-	1
Analysis Date:		01/27/2020	0		01/27/202	0
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores						
Basidiospores						
Chaetomium						
Cladosporium						
Epicoccum						
Ganoderma						
Other brown						
Other colorless						
Penicillium/Aspergillus types†						
Pithomyces						
Rusts						
Scopulariopsis						
Smuts, Periconia, Myxomycetes	1	100	13			
Stachybotrys						
Stemphylium						
Torula						
Ulocladium						
Zygomycetes						
Background debris (1-4+)††	2+			2+		
Hyphal fragments/m3	< 13			< 13		
Pollen/m3	< 13			< 13		
Skin cells (1-4+)	2+			2+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			13			< 13

Comments: A) No spores detected.

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m<sup>3</sup>) multiplied by the sample volume (in liters) divided by 1000 liters.

<sup>†</sup> The spores of Aspergillus and Penicillium (and others such as Acremonium, Paecilomyces) are small and round with very few distinguishing characteristics. They cannot be differentiated by non-viable sampling methods. Also, some species with very small spores are easily missed, and

<sup>††</sup>Background debris indicates the amount of non-biological particulate matter present on the trace (dust in the air) and the resulting visibility for the analyst. It is rated from 1+ (low) to 4+ (high). Counts from areas with 4+ background debris should be regarded as minimal counts and may be higher than reported. It is important to account for samples volumes when evaluating dust levels.

For more information regarding analytical sensitivity, please contact QA by calling the laboratory. ‡ 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".

<sup>§</sup> Total Spores/m3 has been rounded to two significant figures to reflect analytical precision.

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S. South

### SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	E. Stroudsbu	003: 1rg High Sch 302 Scienc	ool South- Rm.	South- Rm. E. Stroudsburg High School So 209		
Comments (see below)		None		None		
Lab ID-Version‡:		11146290-	1		11146291-	1
Analysis Date:		01/27/202	0		01/27/202	0
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores	1	25	53			
Basidiospores	1	25	53	1	25	53
Chaetomium						
Cladosporium						
Epicoccum						
Ganoderma						
Other brown						
Other colorless						
Penicillium/Aspergillus types†	1	25	53	1	25	53
Pithomyces						····
Rusts						
Scopulariopsis						
Smuts, Periconia, Myxomycetes			·	1	100	13
Stachybotrys			Mr-1-10-10			
Stemphylium						
Torula	MANAGEM AND					
Ulocladium						
Zygomycetes			gerandelin versical de la companya d			
Background debris (1-4+)††	2+			2+		
Hyphal fragments/m3	< 13			< 13		
Pollen/m3	< 13			< 13		
Skin cells (1-4+)	1+			2+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			160			120

#### Comments:

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m<sup>3</sup>) multiplied by the sample volume (in liters) divided by 1000 liters.

<sup>†</sup> The spores of Aspergillus and Penicillium (and others such as Acremonium, Paecilomyces) are small and round with very few distinguishing characteristics. They cannot be differentiated by non-viable sampling methods. Also, some species with very small spores are easily missed, and may be undercounted.

<sup>††</sup>Background debris indicates the amount of non-biological particulate matter present on the trace (dust in the air) and the resulting visibility for the analyst. It is rated from 1+ (low) to 4+ (high). Counts from areas with 4+ background debris should be regarded as minimal counts and may be higher than reported. It is important to account for samples volumes when evaluating dust levels.

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S. South

## SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:		005:		006:			
	E. Stroudsburg High School South- Aux.						
		_			26		
	Gym						
Comments (see below)		A			None		
Lab ID-Version‡:		11146292-			11146293-		
Analysis Date:		01/27/2020	)		01/27/202	0	
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3	
Ascospores							
Basidiospores				1	25	53	
Chaetomium							
Cladosporium							
Epicoccum							
Ganoderma							
Other brown							
Penicillium/Aspergillus types†							
Pithomyces							
Rusts							
Scopulariopsis							
Smuts, Periconia, Myxomycetes							
Stachybotrys							
Stemphylium							
Torula							
Ulocladium							
Zygomycetes							
Background debris (1-4+)††	1+			1+			
Hyphal fragments/m3	< 13			< 13			
Pollen/m3	< 13			13			
Skin cells (1-4+)	1+			1+			
Sample volume (liters)	75			75			
§ TOTAL SPORES/m3			< 13			53	

Comments: A) No spores detected.

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m<sup>3</sup>) multiplied by the sample volume (in liters) divided by 1000 liters.

<sup>†</sup> The spores of Aspergillus and Penicillium (and others such as Acremonium, Paecilomyces) are small and round with very few distinguishing characteristics. They cannot be differentiated by non-viable sampling methods. Also, some species with very small spores are easily missed, and may be undercounted.

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S. South

# SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	E. Stroudsbu	007: 1rg High Sch 20	ool South- Rm.	008: E. Stroudsburg High School South- 23		
Comments (see below)	- 112	None		None		
Lab ID-Version‡:		11146294-	1		11146295-	1
Analysis Date:	***************************************	01/27/2020	0		01/27/202	0
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores						
Basidiospores	1	25	53			
Chaetomium						
Cladosporium	1	25	53			
Epicoccum						
Ganoderma						
Other brown						
Other colorless						
Penicillium/Aspergillus types†						
Pithomyces						
Rusts						
Scopulariopsis	2014-21.					
Smuts, Periconia, Myxomycetes				1	100	13
Stachybotrys						
Stemphylium						
Torula						
Ulocladium						
Zygomycetes						
Background debris (1-4+)††	2+			1+		
Hyphal fragments/m3	< 13			13		
Pollen/m3	< 13			13		
Skin cells (1-4+)	1+			1+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			110			13

**Comments:** 

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m<sup>3</sup>) multiplied by the sample volume (in liters) divided by 1000 liters.

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<sup>§</sup> Total Spores/m3 has been rounded to two significant figures to reflect analytical precision.

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S. South

### SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	E. Stroudsbu	009: 1rg High Sch 226	ool South- Rm.	010: E. Stroudsburg High School Sout 118		
Comments (see below)		None		None		
Lab ID-Version‡:		11146296-	1		11146297-	1
Analysis Date:	W.W.	01/27/2020	0		01/27/202	0
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores						
Basidiospores	2	25	110			
Chaetomium						
Cladosporium						
Epicoccum						
Ganoderma						
Other brown						
Other colorless						
Penicillium/Aspergillus types†						
Pithomyces						
Rusts						
Scopulariopsis				1	25	53
Smuts, Periconia, Myxomycetes						
Stachybotrys				1	100	13
Stemphylium						
Torula						
Ulocladium						
Zygomycetes						
Background debris (1-4+)††	1+			1+		
Hyphal fragments/m3	13			< 13		
Pollen/m3	< 13			< 13		
Skin cells (1-4+)	1+		***************************************	1+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			110			67

Comments:

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw

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S. South

# SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	E. Stroudsbu	011: irg High Sch 114	ool South- Rm.	012: E. Stroudsburg High School South- 88		
Comments (see below)		None		None		
Lab ID-Version‡:		11146298-	1		11146299-	1
Analysis Date:	*****	01/27/2020	0		01/27/202	
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores						
Basidiospores	1	25	53	1	25	53
Chaetomium						
Cladosporium						
Epicoccum						
Ganoderma						
Other brown						
Other colorless						
Penicillium/Aspergillus types†	11	25	53			***************************************
Pithomyces						
Rusts						
Scopulariopsis						
Smuts, Periconia, Myxomycetes						
Stachybotrys						
Stemphylium						,,,,,
Torula						
Ulocladium						3000
Zygomycetes						
Background debris (1-4+)††	1+			1+		
Hyphal fragments/m3	< 13			< 13		7.00
Pollen/m3	< 13			< 13		
Skin cells (1-4+)	1+			1+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			110			53

Comments:

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m^3) multiplied by the sample volume (in liters) divided by 1000 liters.

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S. South

## SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	013: E. Stroudsburg High School South- Rm. 92			014: E. Stroudsburg High School South- Rm 101		
Comments (see below)		None		None		
Lab ID-Version‡:		11146300-	1		11146301-	1
Analysis Date:		01/27/202	0		01/27/202	0
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3
Ascospores						
Basidiospores				2	25	110
Chaetomium						
Cladosporium						
Epicoccum	1	100	13			
Ganoderma				1	25	53
Other brown	1	25	53			
Other colorless						
Penicillium/Aspergillus types†						
Pithomyces						
Rusts						
Scopulariopsis						
Smuts, Periconia, Myxomycetes						
Stachybotrys						
Stemphylium	******					
Torula						
Ulocladium						
Zygomycetes						
Background debris (1-4+)††	2+			1+		
Hyphal fragments/m3	< 13			< 13		
Pollen/m3	< 13			< 13		
Skin cells (1-4+)	2+			1+		
Sample volume (liters)	75			75		
§ TOTAL SPORES/m3			67			160

Comments:

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

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S. South

## SPORE TRAP REPORT: NON-VIABLE METHODOLOGY

Location:	E. Stroudsbu	015: E. Stroudsburg High School South- Rm. 106			016: E. Stroudsburg High School Sou Exterior		
Comments (see below)		None		None			
Lab ID-Version‡:		11146302-	1		11146303-	1	
Analysis Date:		01/27/202	0		01/27/202	0	
	raw ct.	% read	spores/m3	raw ct.	% read	spores/m3	
Ascospores							
Basidiospores	2	25	110	4	25	210	
Chaetomium							
Cladosporium							
Epicoccum							
Ganoderma							
Other brown						***************************************	
Other colorless							
Penicillium/Aspergillus types†	1	25	53				
Pithomyces							
Rusts							
Scopulariopsis							
Smuts, Periconia, Myxomycetes							
Stachybotrys							
Stemphylium							
Torula							
Ulocladium	4-4						
Zygomycetes							
Background debris (1-4+)††	2+			2+		,	
Hyphal fragments/m3	13			< 13			
Pollen/m3	< 13			< 13			
Skin cells (1-4+)	2+			< 1+			
Sample volume (liters)	75			75			
§ TOTAL SPORES/m3			160			210	

**Comments:** 

Spore types listed without a count or data entry were not detected during the course of the analysis for the respective sample, indicating a raw count of <1 spore.

The analytical sensitivity is the spores/m<sup>3</sup> divided by the raw count, expressed in spores/m<sup>3</sup>. The limit of detection is the analytical sensitivity (in spores/m<sup>3</sup>) multiplied by the sample volume (in liters) divided by 1000 liters.

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