

HL11-0301

 Sample ID: BIA250728S0014
 Strain: Acapulco Gold

 Produced:
 Collected:
 Received: 07/28/2025
 Completed: 07/31/2025
 Batch#:

 Client
Turkey Hollow Cannabis
 Lic. #
 Pob 192
 S Pomfret, VT 05067

 Matrix: Plant
 Type: Flower - Cured
 Sample Size: 4.98 g
 Lot#:


Summary

Test	Date Tested	Result
Sample		Complete
Cannabinoids	07/30/2025	Complete
Moisture	07/28/2025	10.60% - Complete
Water Activity	07/28/2025	0.534 aw - Complete
Terpenes	07/29/2025	Complete
Microbials	07/31/2025	Complete

Cannabinoids

Completed

28.32%			0.06%			37.07%			
Total THC			Total CBD			Total Cannabinoids			
Analyte	LOQ	Results	Results	Mass	Analyte	LOQ	Results	Results	Mass
	mg/g	%	mg/g	mg/serving		mg/g	%	mg/g	mg/serving
CBDVa	0.0003	<LOQ	<LOQ		CBCVa	0.0003	<LOQ	<LOQ	
CBDV	0.0003	<LOQ	<LOQ		CBNa	0.0003	<LOQ	<LOQ	
CBDa	0.0005	0.07	0.7		Δ9-THC	0.0005	1.01	10.1	
CBGa	0.0005	2.93	29.3		Δ8-THC	0.0003	0.05	0.5	
CBG	0.0005	0.28	2.8		Δ10-THC*	0.0002	<LOQ	<LOQ	
CBD	0.0005	<LOQ	<LOQ		CBL	0.0005	<LOQ	<LOQ	
THCV	0.0003	<LOQ	<LOQ		CBC	0.0003	<LOQ	<LOQ	
CBLV	0.0003	0.07	0.7		THCa	0.0005	31.14	311.4	
CBCV	0.0003	<LOQ	<LOQ		CBCa	0.0006	0.49	4.9	
THCVa	0.0003	1.03	10.3		CBLa	0.0005	<LOQ	<LOQ	
CBN	0.0005	<LOQ	<LOQ		Total THC		28.32	283.19	
					Total CBD		0.06	0.64	
					Total		37.07	370.68	0.00

Analyst: 056

Cannabinoids Methodology: High Performance Liquid Chromatography (HPLC) using PerkinElmer FLEXAR™ with Photo Diode Array Detector (PDA)

Total CBD and total THC are calculated values, to account for assumed decarboxylation from the acid form (THCA or CBDA) to the neutral form, causing weight loss of the acid group. These values are calculated as follows:

$$\text{Total THC} = (\text{THCA} \times 0.877) + \Delta 9\text{-THC}$$

$$\text{Total CBD} = (\text{CBDA} \times 0.877) + \text{CBD Reagent}$$

Blanks: < LOQs for all analytes

LOQ = The lowest quantity that this method can reliably detect. Any cannabinoid that was not detected is assumed to be less than the stated LOQ (<LOQ).

All results reflect dry weight of material, based on % moisture of the sample.

Measurement of Uncertainty (MU): the parameter, associated with the result of a measurement, that characterizes the dispersion of the values that could reasonably be attributed to the

particular quantity subject to measurement. Δ9-THC MU = ±0.005% Total THC MU = ±0.007%

All other cannabinoid MU values are available upon request.

All moisture and water activity analysis is determined by dewpoint measurement using an AQUALAB water activity meter.

*The result is the sum of delta-10 isomers.




 Luke Emerson-Mason
 Laboratory Director
 07/31/2025

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Terpenes

Completed

Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	9.798	0.980
Terpinolene	0.010	8.284	0.828
β-Pinene	0.010	6.479	0.648
Ocimene	0.010	6.422	0.642
3-Carene	0.010	4.956	0.496
β-Myrcene	0.010	4.689	0.469
α-Pinene	0.010	3.309	0.331
α-Terpinene	0.010	1.204	0.120
β-Caryophyllene	0.010	1.168	0.117
γ-Terpinene	0.010	0.822	0.082
Linalool	0.010	0.751	0.075
α-Humulene	0.010	0.592	0.059
trans-Nerolidol	0.010	0.266	0.027
Eucalyptol	0.010	0.162	0.016
Isopulegol	0.010	0.152	0.015
Camphene	0.010	0.125	0.013
Geraniol	0.010	0.096	0.010
Guaiol	0.010	0.075	0.008
Caryophyllene Oxide	0.010	0.010	0.001
α-Bisabolol	0.010	0.010	0.001
cis-Nerolidol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
Total		49.373	4.937

Primary Aromas



Analyst: 052

LOQ = The lowest quantity this method can reliably detect. Any terpene that was not detected is assumed to be less than the stated LOQ (<LOQ).

Terpene Methodology: Headspace Sampler, Gas Chromatography-Mass Spectrometry (GC-MS), using Perkin Elmer Clarus® SQ8 GC MS

Reagent Blanks: < LOQs for all analytes

All results reflect dry weight of material, based on % moisture of the sample.

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Pathogens

Completed

Pathogens	LOD	Results
	CFU/g	CFU/g
Aspergillus	5	Not Detected
Shiga Toxin E. Coli	5	Not Detected
Salmonella SPP	5	Not Detected

Analyst: 018

Test Methodology: Bio-Rad IQ-Check PCR Kits

cfu/g = colony forming units per gram

LOD = The lowest quantity that this method can reliably detect. Any microbial growth that was not detected is assumed to be less than the stated LOD (<LOD).

Reagent Blanks: <LOD for all analytes




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