

0203-15-01

 Sample ID: BIA250724S0004
 Strain: Garlic crusher

 Matrix: Plant
 Type: Flower - Cured
 Sample Size: 6.65 g
 Lot#:

 Produced:
 Collected:
 Received: 07/24/2025
 Completed: 07/31/2025
 Batch#:

 Client
cloud 9
 Lic. #
 4082 Noyestar Rd
 East Hardwick, VT 05836


Summary

Test	Date Tested	Result
Sample		Complete
Cannabinoids	07/28/2025	Complete
Moisture	07/25/2025	11.20% - Complete
Water Activity	07/25/2025	0.560 aw - Complete
Terpenes	07/24/2025	Complete
Microbials	07/31/2025	Complete

Cannabinoids

Completed

29.88%

Total THC

0.11%

Total CBD

36.35%

Total Cannabinoids

Analyte	LOQ	Results	Results	Mass
	mg/g	%	mg/g	mg/serving
CBDVa	0.0003	<LOQ	<LOQ	
CBDV	0.0003	<LOQ	<LOQ	
CBDa	0.0005	0.12	1.2	
CBGa	0.0005	0.94	9.4	
CBG	0.0005	0.15	1.5	
CBD	0.0005	<LOQ	<LOQ	
THCV	0.0003	<LOQ	<LOQ	
CBLV	0.0003	<LOQ	<LOQ	
CBCV	0.0003	0.05	0.5	
THCVA	0.0003	0.80	8.0	
CBN	0.0005	<LOQ	<LOQ	

Analyte	LOQ	Results	Results	Mass
	mg/g	%	mg/g	mg/serving
CBCVa	0.0003	<LOQ	<LOQ	
CBNa	0.0003	<LOQ	<LOQ	
Δ9-THC	0.0005	0.15	1.5	
Δ8-THC	0.0003	<LOQ	<LOQ	
Δ10-THC*	0.0002	<LOQ	<LOQ	
CBL	0.0005	<LOQ	<LOQ	
CBC	0.0003	<LOQ	<LOQ	
THCa	0.0005	33.90	339.0	
CBCa	0.0006	0.25	2.5	
CBLa	0.0005	<LOQ	<LOQ	
Total THC		29.88	298.83	
Total CBD		0.11	1.08	
Total		36.35	363.54	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:

TotalTHC=(THCAx0.877)+Δ9-THC

Total CBD = (CBDA x 0.877) + 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).

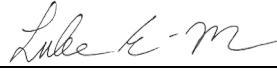
All results reflect dry weight of material, based on % moisture of the sample.

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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 (866) 506-5866
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Terpenes

Completed

Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	5.809	0.581
Ocimene	0.010	4.144	0.414
β-Myrcene	0.010	2.833	0.283
β-Caryophyllene	0.010	2.470	0.247
Linalool	0.010	1.638	0.164
β-Pinene	0.010	1.425	0.142
α-Humulene	0.010	1.130	0.113
α-Pinene	0.010	0.880	0.088
Isopulegol	0.010	0.522	0.052
Terpinolene	0.010	0.183	0.018
Camphehe	0.010	0.172	0.017
α-Bisabolol	0.010	0.030	0.003
γ-Terpinene	0.010	0.014	0.001
3-Carene	0.010	<LOQ	<LOQ
α-Terpinene	0.010	<LOQ	<LOQ
Caryophyllene Oxide	0.010	<LOQ	<LOQ
cis-Nerolidol	0.010	<LOQ	<LOQ
Eucalyptol	0.010	<LOQ	<LOQ
Geraniol	0.010	<LOQ	<LOQ
Guaiol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
trans-Nerolidol	0.010	<LOQ	<LOQ
Total		21.252	2.125

Primary Aromas



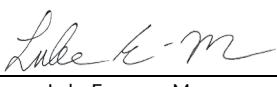
Analyst: 052

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.

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



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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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