

scLT 0203-12-02

 Sample ID: BIA241011S0016
 Strain: golden goat

 Matrix: Plant
 Type: Flower - Cured
 Sample Size: 6.46 g
 Lot#:

 Produced:
 Collected:
 Received: 10/11/2024
 Completed: 10/17/2024
 Batch#:

 Client
cloud 9
 Lic. # scLT0203
 4082 Noyestar Rd
 East Hardwick, VT 05836


Summary

Test	Date Tested	Result
Sample	10/15/2024	Complete
Cannabinoids	10/14/2024	Complete
Moisture	10/14/2024	11.30% - Complete
Water Activity	10/14/2024	0.567 aw - Complete
Terpenes	10/14/2024	Complete
Microbials	10/17/2024	Complete

Cannabinoids

Completed

27.93%	0.07%	32.55%
Total THC	Total CBD	Total Cannabinoids

Analyte	LOQ	Results	Results	Mass
	mg/g	%	mg/g	mg/serving
CBDVa	0.0005	<LOQ	<LOQ	
CBDV	0.0012	<LOQ	<LOQ	
CBDa	0.0008	0.08	0.8	
CBGa	0.0008	0.68	6.8	
CBG	0.0019	0.09	0.9	
CBD	0.0019	<LOQ	<LOQ	
THCV	0.0021	<LOQ	<LOQ	
CBN	0.0013	<LOQ	<LOQ	
Δ9-THC	0.0020	0.99	9.9	
Δ8-THC	0.0019	<LOQ	<LOQ	
Δ10-THC	0.0002	<LOQ	<LOQ	
CBC	0.0024	<LOQ	<LOQ	
THCa	0.0034	30.72	307.2	
Total THC		27.93	279.29	
Total CBD		0.07	0.67	
Total		32.55	325.53	0.00

Analyst: 048

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

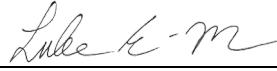
LOQ = The lowest quantity that this method can reliably detect. Any cannabinoid that was not detected is assumed to be less than the stated LOQ (<LOQ).

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. $\Delta 9\text{-THC MU} = \pm 0.005\%$ $\text{Total THC MU} = \pm 0.007\%$

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.

 Luke Emerson-Mason
 Laboratory Director
 10/17/2024

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 (866) 506-5866
www.confidentlims.com


sclt 0203-12-02

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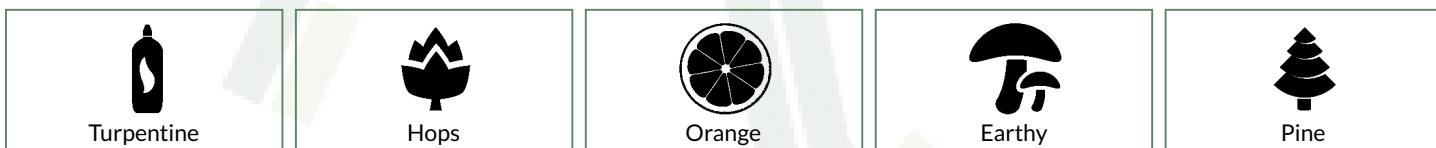
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Terpenes

Completed

Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Terpinolene	0.010	4.415	0.441
β-Myrcene	0.010	3.786	0.379
Limonene	0.010	3.614	0.361
Ocimene	0.010	3.315	0.331
α-Pinene	0.010	2.557	0.256
β-Pinene	0.010	2.164	0.216
β-Caryophyllene	0.010	1.617	0.162
Linalool	0.010	0.935	0.093
3-Carene	0.010	0.823	0.082
α-Humulene	0.010	0.693	0.069
α-Terpinene	0.010	0.304	0.030
γ-Terpinene	0.010	0.272	0.027
Guaiol	0.010	0.152	0.015
Eucalyptol	0.010	0.115	0.012
Camphene	0.010	0.054	0.005
α-Bisabolol	0.010	0.033	0.003
Caryophyllene Oxide	0.010	<LOQ	<LOQ
cis-Nerolidol	0.010	<LOQ	<LOQ
Geraniol	0.010	<LOQ	<LOQ
Isopulegol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
trans-Nerolidol	0.010	<LOQ	<LOQ
Total		24.848	2.485

Primary Aromas


Analyst: 045

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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 10/17/2024

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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

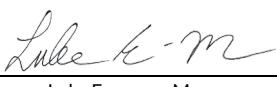
Analyst: 018

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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