

Melonhead mother

Sample ID: BIA240508S0004 Strain: Melonhead

Matrix: Plant Type: Flower - Cured Sample Size: 5 g Lot#:

Produced: Collected: Received: 05/14/2024 Completed: 05/23/2024

Nightime land productions Lic. # SCLT0366 646 Rte 100N Wilmington, VT 05363



Summary

Test	Date Tested	Result
Sample		In Progress
Cannabinoids	05/20/2024	Complete
Moisture	05/15/2024	11.60% - Complete
Water Activity	05/15/2024	0.580 aw - Complete
Terpenes	05/16/2024	Complete
Microbials	05/23/2024	Complete
Pesticides	05/17/2024	Complete

Cannabinoids Completed

1	19.65% otal THC		0.06% Total CBD		23.41% Total Cannabinoids
Analyte	LOQ	Results	Results	Mass	
CBDVa CBDV CBDa CBGa CBG CBD THCV	mg/g 0.0005 0.0012 0.0008 0.0008 0.0019 0.0019 0.0021	% <loq <loq 0.07 1.01 0.08 <loq <loq< td=""><td>mg/g <loq <loq 0.7 10.1 0.8 <loq <loq< td=""><td>mg/serving</td><td></td></loq<></loq </loq </loq </td></loq<></loq </loq </loq 	mg/g <loq <loq 0.7 10.1 0.8 <loq <loq< td=""><td>mg/serving</td><td></td></loq<></loq </loq </loq 	mg/serving	
CBN Δ9-THC Δ8-THC THCa CBC Total THC Total CBD Total	0.0013 0.0020 0.0019 0.0034 0.0024	<loq 1.08 <loq 21.17 <loq 19.65 0.06 23.41</loq </loq </loq 	<loq 10.8 <loq 211.7 <loq 196.46 0.61 234.12</loq </loq </loq 	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. $\Delta 9$ -THC MU = $\pm 0.005\%$ Total THC MU = $\pm 0.007\%$ All other cannabinoid MU values are available upon request.

All moisture analysis is determined by loss-on-drying measurement using OHAUS Model MB90 Moisture Content Readers.



Luke Emerson-Mason

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Bia Diagnostics Colchester, VT 05446

(802) 540-0148 480 Hercules Drive Suite 101 https://www.biadiagnostics.com/ Lic#TLAB0029

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In Progress Terpenes

	100	.	. .
Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	0.590	0.059
Linalool	0.010	0.491	0.049
Ocimene	0.010	0.343	0.034
β-Caryophyllene	0.010	0.268	0.027
β-Myrcene	0.010	0.264	0.026
β-Pinene	0.010	0.150	0.015
α-H <mark>umulene</mark>	0.010	0.094	0.009
α-Pinene	0.010	0.080	0.008
Geraniol	0.010	0.044	0.004
Guaiol	0.010	0.025	0.002
Camphene	0.010	0.020	0.002
3-Carene	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
α-Bisabolol	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
α-Terpinene	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
Caryophyllene Oxide	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
cis-Nerolidol	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
Eucalyptol	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
y-Terpinene	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
Isopulegol	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
p-Cymene	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
Terpinolene	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
trans-Nerolidol	0.010	<loq< td=""><td><loq< td=""></loq<></td></loq<>	<loq< td=""></loq<>
Total		2.369	0.237
Aromas			

Primary Aromas











Analyst: 048

LOQ = The lowest quantity this method can reliably detect. Any terpene that was not detected is assumed to be less than the stated 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 analysis is determined by loss-on-drying measurement using OHAUS Model MB90 Moisture Content Readers.



Luke Emerson-Mason Laboratory Director

05/23/2024

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Completed **Pesticides**

Category 1 Pesticides	LOQ	Results
	PPM	PPM
Chlorpyrifos	0.0010	<loq< td=""></loq<>
Imazalil	0.0010	<loq< th=""></loq<>
Category 2 Pesticides	LOQ	Results
	PPM	PPM
Abamectin	0.0100	<loq< td=""></loq<>
Acephate	0.0010	<loq< td=""></loq<>
Acequinocyl	0.0010	<loq< td=""></loq<>
Azoxystrobin	0.0010	<loq< td=""></loq<>
Bifenazate	0.0010	<loq< td=""></loq<>
Bifenthrin	0.0010	<loq< td=""></loq<>
Carbaryl	0.0010	<loq< td=""></loq<>
Cypermethrin	0.0100	<loq< td=""></loq<>
Etoxazole	0.0010	<loq< td=""></loq<>
Imidacloprid	0.0010	<loq< td=""></loq<>
Myclobutanil	0.0010	<loq< td=""></loq<>
Spinosyn A	0.0010	<loq< td=""></loq<>
Spinosyn D	0.0010	<loq< td=""></loq<>

Analyst: 045

Pesticides Methodology: Liquid Chromatography with Tandem Mass Spectrometry using PerkinElme QSight® LX50 UHPLC and QSight 220 Mass Spectrometer

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

ppm = parts per million

All moisture analysis is determined by loss-on-drying measurement using OHAUS Model MB90 Moisture Content Readers.



Luke Emerson-Mason Laboratory Director 05/23/2024

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Bia Diagnostics

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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: 049

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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