

6.06

Sample ID: BIA241104S0005 Strain: Cherry Poison

Matrix: Plant Type: Flower - Cured Sample Size: 8.76 g Lot#: Produced: Collected: Received: 11/04/2024 Completed: 11/11/2024 Batch#:

**Bia Diagnostics** 

Colchester, VT 05446

480 Hercules Drive Suite 101

(802) 540-0148 https://www.biadiagnostics.com/ Lic# TLAB0029

Client

Clovis LLC

Lic. # CLTV0099

506 Marcoux Rd

Hyde Park, VT 05655

QA Testing

1 of 3

Summary Test Date Tested Result Sample Complete Cannabinoids 11/06/2024 Complete 6.06 Cherry Pois CLOVIS CLTV0099-006 Moisture 11/05/2024 9.50% - Complete Water Activity 11/05/2024 0.464 aw - Complete Terpenes 11/06/2024 Complete 11/08/2024 Microbials Complete

## Cannabinoids

<b>19.24%</b> Total THC		0.07% Total CBD			<b>23.53%</b> Total Cannabinoids
Analyte	LOQ	Results	Results	Mass	
CBDVa CBDV CBDa CBGa CBG CBD THCV CBN A9-THC A8-THC A10-THC CBC THCa Total THC	mg/g 0.0005 0.0012 0.0008 0.0019 0.0019 0.0019 0.0021 0.0013 0.0020 0.0019 0.0020 0.0019 0.0002 0.0002	% <loq 0.08 1.41 0.11 <loq <loq <loq 0.31 <loq 0.04 <loq 21.59 <b>19.24</b></loq </loq </loq </loq </loq </loq 	mg/g <loq <loq 0.8 14.1 1.1 <loq <loq <loq 3.1 <loq 0.4 <loq 0.4 <loq 215.9 <b>192.43</b></loq </loq </loq </loq </loq </loq </loq </loq 	mg/serving	
Total CBD Total		0.07 23.53	0.66 235.32	0.00	

Analyst: 056

Cannabinoids Methodology: High Performance Liquid Chromatography (HPLC) using PerkinElmer FLEXAR TM 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 = ±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.



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Luke Emerson-Mason

Laboratory Director 11/11/2024 Confident LIMS All Rights Reserved coa.support@confidentlims.com (866) 506-5866 www.confidentlims.com



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Completed



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## Terpenes

Completed LOC Results Results Analyte mg/g mg/g Limonene 0.010 6.380 0.638 0.010 2.856 0.286 Terpinolene 0.010 2.558 0.256 Ocimene β-Myrcene 0.010 2.462 0.246 β-Caryophyllene 0.010 2.441 0.244 0.235 α-Pinene 0.010 2.354 **β-Pinene** 2.204 0.220 0.010 Linalool 1.216 0.122 0.010 α-Humulene 0.010 0.967 0.097 3-Carene 0.010 0.352 0.035 Camphene 0.010 0.170 0.017 α-Terpinene 0.010 0.155 0.015 Eucalyptol 0.010 0.151 0.015 v-Terpinene 0.0100.110 0.011 0.070 Geraniol 0.010 0.007 0.010 0.040 0.004 Guaiol Caryophyllene Oxide 0.010 0.020 0.002 α-Bisabolol 0.010 <LOO <LOQ cis-Nerolidol 0.010 <LOQ <LOQ Isopulegol 0.010 <LOQ <LOQ p-Cymene 0.010 <LOQ <LOQ trans-Nerolidol 0.010 <LOQ <LOQ

Total Primary Aromas

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Orange	Turpentine	Earthy	Hops	Cinnamon

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 (<LOQ).

Terpene Methodology: Headspace Sampler, Gas Chromatography-Mass Spectrometry (GC-MS), using Perkin Elmer Clarus® SQ8 GC MS Reagent Blanks: < LOQs for all analytes

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MW C Luke Emerson-Mason

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2.450

24.505



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## Pathogens

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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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