

Big Perm

Sample ID: BIA251126S0779
 Strain: Manu0114-bigperm
 Harvest Lot:
 Matrix: Concentrates & Extracts
 Type: Rosin
 Sample Size: 1 units
 Lot#:

Produced:
 Collected:
 Received: 11/26/2025
 Completed: 12/05/2025
 Batch#:

Client
 PermaGanix



Summary

Test	Date Tested	Result
Sample		Complete
Cannabinoids	12/02/2025	Complete
Terpenes	12/03/2025	Complete
Microbials	12/05/2025	Complete

Cannabinoids

Completed

66.73%

Total THC

ND

Total CBD

83.46%

Total Cannabinoids

Analyte	LOQ	Results	Results	Mass	Mass
	%	%	mg/g	mg/mL	mg/container
CBDVa	0.0000	<LOQ	<LOQ		
CBDV	0.0000	<LOQ	<LOQ		
CBDa	0.0001	<LOQ	<LOQ		
CBGa	0.0001	4.76	47.6		
CBG	0.0001	0.41	4.1		
CBD	0.0001	<LOQ	<LOQ		
THCV	0.0000	<LOQ	<LOQ		
CBLV	0.0000	0.27	2.7		
CBCV	0.0000	<LOQ	<LOQ		
THCVA	0.0000	0.39	3.9		
CBN	0.0001	<LOQ	<LOQ		

Analyte	LOQ	Results	Results	Mass	Mass
	%	%	mg/g	mg/mL	mg/container
CBCVa	0.0000	<LOQ	<LOQ		
CBNa	0.0000	<LOQ	<LOQ		
Δ9-THC	0.0001	1.39	13.9		
Δ8-THC	0.0000	<LOQ	<LOQ		
Δ10-THC*	0.0000	<LOQ	<LOQ		
CBL	0.0001	<LOQ	<LOQ		
CBC	0.0000	<LOQ	<LOQ		
THCa	0.0001	74.49	744.9		
CBCa	0.0001	1.74	17.4		
CBLa	0.0001	<LOQ	<LOQ		
Total THC		66.73	667.26		
Total CBD		ND	ND	ND	ND
Total		83.46	834.55	0.00	0.00

Analyst: 052

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:

Total THC = (THCA x 0.877) + 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
 12/05/2025

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 (866) 506-5866
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Terpenes

Completed

Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	3.837	0.384
Ocimene	0.010	2.872	0.287
Linalool	0.010	2.618	0.262
β-Myrcene	0.010	2.423	0.242
β-Caryophyllene	0.010	2.420	0.242
β-Pinene	0.010	2.289	0.229
α-Pinene	0.010	1.899	0.190
α-Humulene	0.010	0.932	0.093
Terpinolene	0.010	0.415	0.042
Camphene	0.010	0.354	0.035
α-Bisabolol	0.010	0.046	0.005
γ-Terpinene	0.010	0.041	0.004
Geraniol	0.010	0.028	0.003
Guaiol	0.010	0.027	0.003
α-Terpinene	0.010	0.022	0.002
3-Carene	0.010	0.015	0.002
Caryophyllene Oxide	0.010	<LOQ	<LOQ
cis-Nerolidol	0.010	<LOQ	<LOQ
Eucalyptol	0.010	<LOQ	<LOQ
Isopulegol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
trans-Nerolidol	0.010	<LOQ	<LOQ
Total		20.238	2.024

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: 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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