Bia Diagnostics Laboratories

Bia Diagnostics
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Lic# TLAB0029

GMO HL17

Sample ID: BIA250807S0011
Strain: GMO

Matrix: Plant
Type: Flower - Cured
Sample Size: 8.16 g
Lot#: HL17

Produced:
Collected:
Received: 08/07/2025
Completed: 08/14/2025
Batch#:

Client
PermaGanix

Summary

Test	Date Tested	Result
Sample		Complete
Cannabinoids	08/11/2025	Complete
Moisture	08/08/2025	11.70% - Complete
Water Activity	08/08/2025	0.583 aw - Complete
Terpenes	08/12/2025	Complete
Microbials	08/14/2025	Complete

Cannabinoids

Completed

32.79%	0.09%	40.67%
Total THC	Total CBD	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	0.10	1.0	
CBGa	0.0005	1.83	18.3	
CBG	0.0005	0.11	1.1	
CBD	0.0005	<LOQ	<LOQ	
THCV	0.0003	0.07	0.7	
CBLV	0.0003	0.14	1.4	
CBCV	0.0003	<LOQ	<LOQ	
THCVa	0.0003	0.21	2.1	
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	0.18	1.8	
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	37.18	371.8	
CBCa	0.0006	0.43	4.3	
CBLa	0.0005	0.42	4.2	
Total THC		**32.79**	**327.90**	
Total CBD		**0.09**	**0.90**	
Total		**40.67**	**406.67**	**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:
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
08/14/2025

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Terpenes

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Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	8.554	0.855
β-Myrcene	0.010	5.012	0.501
Ocimene	0.010	4.414	0.441
β-Caryophyllene	0.010	2.588	0.259
β-Pinene	0.010	2.191	0.219
Linalool	0.010	1.480	0.148
α-Pinene	0.010	1.245	0.124
α-Humulene	0.010	1.179	0.118
Terpinolene	0.010	0.255	0.025
Camphene	0.010	0.174	0.017
α-Bisabolol	0.010	0.021	0.002
γ-Terpinene	0.010	0.020	0.002
α-Terpinene	0.010	0.017	0.002
3-Carene	0.010	0.013	0.001
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
Isopulegol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
trans-Nerolidol	0.010	<LOQ	<LOQ
Total		**27.162**	**2.716**

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 (<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

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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: 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

Luke Emerson-Mason
Laboratory Director
08/14/2025