

Certificate of Analysis

Powered by Confident Cannabis

Groove Solventless

Sample: 2311FID5812.37323

Strain: Grape Cream Cake

Batch #: C; Lot #:

METRC Batch: 1A4080100013EC1000000971; METRC Sample: 1A4080100013EC1000000977 Analysis Initiated: 11/10/2023; Report Created: 11/15/2023

Sampling SOP: SOP-0050

Grape Cream Cake Liquid Flower (HB37.1)

Ingestible, Tincture, Other Harvest/Production Date: 09/21/2023





102.94 mg/unit

Total Potential Psychoactive THC

102.94 mg/unit

Total Raw THC

0.67 mg/unit

Total CBD

111.65 mg/unit

Total Cannabinoids

Not Tested

Foreign Matter

Not Tested

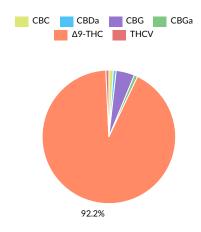
NT

Moisture

Pass Cannabinoids

Analytical Calibration Batch: Cannabinoids AF 10302023

Analyte	LOQ	Mass	Mass
	mg/unit	mg/unit	mg/g
THCa	0.01	ND	ND
Δ9-ΤΗС	0.01	102.94	17.10
Δ8-ΤΗС	0.01	ND	ND
THCVa	0.01	ND	ND
THCV	0.01	0.81	0.14
CBDa	0.01	0.76	0.13
CBD	0.01	ND	ND
CBN	0.01	ND	ND
CBGa	0.01	0.94	0.16
CBG	0.01	4.99	0.83
CBC	0.01	1.20	0.20
Δ10-ΤΗС	0.01	ND	ND
Total		111.65	18.55



1 Unit = 1 Bottle, 6.0190g

Total Potential Psychoactive THC = THCa * 0.877 + d9-THC | Total CBD = CBDa * 0.877 + CBD

LOQ = Limit of Quantitation; NT = Not Tested; NR = Not Reported; ND = None Detected; PPM = Parts per Million; PPB = Parts per Billion; Unless otherwise stated all quality control samples performed within specifications established by the Laboratory.

SOP-0037; Full spectrum cannabinoid analysis by High Performance Liquid Chromatography with UV detection (HPLC-UV). Reported result is based on sample dry weight. SOP-0035; Foreign matter inspection includes but is not limited to hair, insects, stems, and feces. Filth is inspected using a M16-209 stereoscope. Stem measurements are performed using fisher caliners

SOP-0036; Moisture analysis is performed using a Shimadzu moisture analyzer MOC63u UL. Testing listed above may not be included in our current PJLA scope of accreditation. Please consult PJLA Certificate# 102722 for a list of current accredited tests



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Andre Umansky Laboratory Director

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Accredited to ISO/IEC 17025:2017 Accreditation #: 102722

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Terpenes

Analytical Calibration Batch: Terpenes 10122023

Analyte	LOQ	Mass	Mass
	%	%	mg/g
α-Bisabolol	0.008	ND	ND
α-Humulene	0.008	ND	ND
α-Pinene	0.008	ND	ND
α-Terpinene	0.008	ND	ND
β-Caryophyllene	0.008	ND	ND
β-Myrcene	0.008	ND	ND
β-Pinene	0.008	ND	ND
Camphene	0.008	ND	ND
Caryophyllene Oxide	0.008	ND	ND
cis-Nerolidol	0.008	ND	ND
δ-3-Carene	0.008	ND	ND
δ-Limonene	0.008	ND	ND
Eucalyptol	0.008	ND	ND
y-Terpinene	0.008	ND	ND
Geraniol	0.008	ND	ND
Guaiol	0.008	ND	ND
Isopulegol	0.008	ND	ND
Linalool	0.008	ND	ND
Ocimene	0.008	ND	ND
p-Cymene	0.008	ND	ND
Terpinolene	0.008	ND	ND
trans-Nerolidol	0.008	ND	ND
Total		0.000	0

Primary Aromas



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Residual Solvents Not Tested

Analytical Calibration Batch:

Analyte LOQ State Limits Mass Status Analyte LOQ State Limits Mass Status

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SOP-0056; A wide spectrum analysis of Residual Solvents using Gas Chromatography Mass Spectrometry (Thermo Scientific ISQ7000 GCMS).

Mycotoxins	Not lested
Analytical Calibration Batch:	

Analyte LOQ State Limit Mass Status

Microbials	Pa		Pass
Amalista	Lineia	14	Chahira

Allalyte	LIIIII	IVIASS	Jiaius
	CFU/g	CFU/g	
Mold	10000	NR	NT
Aspergillus flavus	Not Detected in 1.0g	ND	Pass
Aspergillus fumigatus	Not Detected in 1.0g	ND	Pass
Aspergillus niger	Not Detected in 1.0g	ND	Pass
Aspergillus terreus	Not Detected in 1.0g	ND	Pass
Salmonella	Not Detected in 1.0g	ND	Pass
STEC	Not Detected in 1.0g	ND	Pass

LOQ = Limit of Quantitation; The reported result is based on a sample weight with the applicable moisture content for that sample; Unless otherwise stated all quality control samples performed within specifications established by the Laboratory. SOP-0048; Mycotoxin screening is performed using Sciex 6500+ LCMSMS with Exion XR front HPLC.

 $LOQ = Limit of Quantitation; TNTC = Too \ Numerous to Count; Unless otherwise stated all quality control samples performed within specifications established by the Laboratory. SOP-0057: STEC and Salmonela analysis on AriaDX qPCR using Medicinal Genomics validated methods. SOP-0063: Aspergillus species specific analysis on AriaDX qPCR using Medicinal Genomics validated methods. SOP-0061: Mold enumeration using Hardy Diagnostics media.$



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