

Certificate of Analysis

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Groove Solventless

Sample: 2502FID0755.4039

Strain: Rainbow Sherbert #11/Mai Tai Batch #: a; Lot #: METRC Batch: 1A4080100013EC1000008358; METRC Sample: 1A4080100013EC1000008365 Analysis Initiated: 02/11/2025; Report Created: 02/17/2025 Sampling SOP: SOP-0050

Rainbow Sherbert #11 (HB84) Flower Infused w/ Mai Tai Solventless Hash (HB87)

Concentrates & Extracts, Infused/Enhanced Preroll, Ice/Water Harvest/Production Date: 02/05/2025

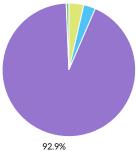
ND Total CBD	Pass Foreign Matter	
43.45%	Pass 8.98% Moisture	
	Total CBD	

Cannabinoids

Analytical Calibration Batch: Cannabinoids AF 01022025

Analyte	LOQ	Mass	Mass
	%	%	mg/g
THCa	0.09	40.35	403.5
Δ9-THC	0.09	1.25	12.5
THCVa	0.09	0.28	2.8
THCV	0.09	ND	ND
CBDa	0.09	ND	ND
CBD	0.09	ND	ND
CBN	0.09	ND	ND
CBGa	0.09	1.57	15.7
CBG	0.09	ND	ND
CBC	0.09	ND	ND
Total		43.45	434.5

CBGa Δ9-THC THCa THCVa



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. 50P-0041: 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, seeds, and feces. SOP-0036: Moisture analysis. 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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Fidelity Diagnostics Laboratory





Andre Umansky Laboratory Director

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

Testing performed onsite at Fidelity Diagnostics using valid testing methodologies and a quality system as required by state law. Fidelity Diagnostics makes no claims as to the efficacy, safety or other risks associated with any detected or non-detected levels of any compounds reported herein. Results valid only for item tested. Unless otherwise stated all quality control samples performed within specifications established by the Laboratory. Certificate shall not be reproduced, without written approval by the laboratory. Quality and uncertainty data available upon request. Sample conformity is set by the state of Montana. Version No. 001

Pass



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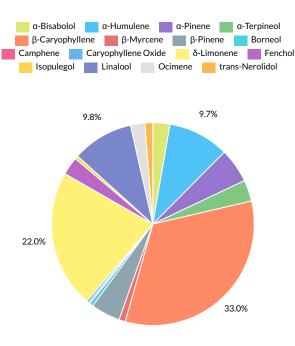
Terpenes

Analyte	LOQ	Result	Result
	%	%	mg/g
β-Caryophyllene	0.008	1.353	13.53
δ-Limonene	0.008	0.901	9.01
Linalool	0.008	0.399	3.99
α-Humulene	0.008	0.399	3.99
α-Pinene	0.008	0.224	2.24
β-Pinene	0.008	0.191	1.91
α-Terpineol	0.005	0.139	1.39
Fenchol	0.003	0.117	1.17
α-Bisabolol	0.008	0.112	1.12
Ocimene	0.008	0.093	0.93
trans-Nerolidol	0.008	0.052	0.52
β-Myrcene	0.008	0.042	0.42
Borneol	0.003	0.024	0.24
Isopulegol	0.008	0.022	0.22
Caryophyllene Oxide	0.008	0.015	0.15
Camphene	0.008	0.010	0.10
α-Phellandrene	0.003	ND	ND
α-Terpinene	0.008	ND	ND
cis-Nerolidol	0.008	ND	ND
δ-3-Carene	0.008	ND	ND
Eucalyptol	0.008	ND	ND
y-Terpinene	0.008	ND	ND
Geraniol	0.008	ND	ND
Geranyl Acetate	0.003	ND	ND
Guaiol	0.008	ND	ND
p-Cymene	0.008	ND	ND
Terpinolene	0.008	ND	ND
Total		4.093	40.93



Analytical Calibration Batch: Terpenes 12/23/2025





Primary Aromas



LOQ = Limit of Quantitation; NT = Not Tested; NR = Not Reported; ND = None Detected. Results based on dry weight. SOP-0037; GC-FID Analysis of Terpenes in Cannabis and Cannabis Products. 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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LOQ = Limit of Quantitation; NT = Not Tested; NR = Not Reported; ND = None Detected. PPM = Parts per Million. SOP-0060; A mass spectrum analysis of Residual Solvents using Gas Chromatography Mass Spectrometry (Thermo Scientific ISQ7000 GCMS).

Mycotoxins		Not Te	ested	Microbi	als		Not T	ested
Analytical Calibration Batch: Analyte	LOO State Limit	Mass	Status	Analyte		Limit	Mass	<u>Status</u>
LOQ = Limit of Quantitation; NT = Not PPB = Parts per Billion. SOP-0042; Pes Sciex 6500+ LCMSMS with Exion XR fr	Tested; NR = Not Reported	1; ND = None [Detected.	Count; SOP-005 Genomics valida	; NR = Not Reported; ND 8: STEC and Salmonela ar ted methods. SOP-0063: icinal Genomics validatec cs media.	nalysis on AriaE Aspergillus spe	X qPCR using Medicina cies specific analysis on	al AriaDX
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