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Sample: 2403TLL0094.0487

Genesis Bioceuticals, LLC

1120 W Watkins St Phoenix, AZ 85007 shonae.j@genbioaz.com (847) 682-4899 Lic. #00000058DCQU00115543 Harvest Dates: 03/06/2024 Mac 1 (Caps Cut) Flower/Pre-Roll

Plant, Flower - Cured

Strain: Mac 1 (Caps Cut) Parent Batch #: ; Batch#: MAC1-2373-20240306; Batch Size: 11 g Sample Received: 03/18/2024; Report Created: 03/25/2024; Expires: 03/25/2025 Manufacturing Date: Sampling: ; Environment:



Dispensary License #: ; Manufacturing License #: ; Cultivation License #:

Cannabinoids

	_		
<loq< th=""><th></th><th>29.31</th><th>%</th></loq<>		29.31	%
Total CBD		Total Canna Q3	binoids
LOQ	Mass	Mass	Qualifier
%	%	mg/g	
0.10	28.21	282.1	
0.10	0.16	1.6	
0.10	ND	ND	
0.10	ND	ND	
0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
0.10	ND	ND	
0.10	ND	ND	
0.10	ND	ND	
0.10	0.95	9.5	
0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
0.10	ND	ND	
	29.31	293.1	
	Total CBD <u>LOQ</u> % 0.10 0.1	Log Mass % % 0.10 28.21 0.10 28.21 0.10 0.16 0.10 ND 0.10 State 0.10 ND 0.10 ND	Total CBD Total Canna Q3 LOQ Mass % % % % % 28.21 0.10 28.21 0.10 0.16 0.10 ND 0.10 ND

Total THC = THCa * $0.877 + \Delta 9$ -THC Total CBD = CBDa * 0.877 + CBDInstrument: HPLC-DAD: ; Method: TPL_Potency_01



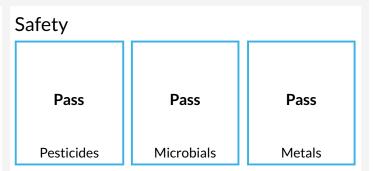
1721 E McDowell Road Phoenix, AZ (602) 368-4233 transparentlabsaz.com Lic# 0000029LRCXG19240160

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



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Terpenes TPL Terpenes 01

lerpenes_01	7		ý	Ĭ
Lemon	Earthy		Ho	ps
Analyte	LOQ	Mass	Mass	Qualifier
δ-Limonene Ocimene α-Humulene β-Caryophyllene α-Pinene Linalool trans-Nerolidol β-Pinene β-Myrcene Terpinolene y-Terpinene Eucalyptol Camphene 3-Carene α-Bisabolol α-Terpinene Caryophyllene Oxide cis-Nerolidol	%	% 0.4300 0.3900 0.3300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2300 0.2000 0.2000 0.2000 0.4000 0.2000 0.2000 0.2000 0.2000 0.2000 0.2000 0.2300 0.200000000	mg/g 4.300 3.900 3.300 2.300 2.300 2.300 2.300 2.300 2.300 0.200 0.400 0.200 < < < < < <	33333333333333333333333333333333333333
Geraniol Guaiol		< < <	< < <	Q3 Q3 Q3 Q3 Q3
Isopulegol p-Cymene Total		2.6900	ج 26.900	Q3 Q3
Instrument: CCMS: Method	TDI Town 01			

Instrument: GCMS; Method: TPL_Terp_01 Notes:



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Sampling: ; Environment:



Pass

Plant, Flower - Cured

Dispensary License #: ; Manufacturing License #: ; Cultivation License #:

Pesticides TPL_Pesticides_01

Analyte	LOQ	Limit	Mass	Status C	Qualifier	Analyte	LOQ	Limit	Mass	Status C	Qualifier
	PPM	PPM	PPM		<u> </u>		PPM	PPM	PPM		<u> </u>
A have a ativ	0.24	0.50		Deee	L1 R2	Hexythiazox	0.48	1.00	ND	Pass	
Abamectin	0.24	0.50	ND	Pass	M1	Imazalil	0.10	0.20	ND	Pass	
Acephate	0.19	0.40	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	M2
Aldicarb	0.19	0.40	ND	Pass		Methyl	0.19	0.40	ND	PdSS	IVIZ
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenazate	0.10	0.20	ND	Pass	M2	Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass		Methomyl	0.19	0.40	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	
Carbofuran	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass	R1 M2	Oxamyl	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass		Paclobutrazol	0.19	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2
Clofentezine	0.10	0.20	ND	Pass	M1	Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass	M1	Piperonyl	0.96	2.00	<loq< th=""><th>Pass</th><th></th></loq<>	Pass	
Cypermethrin	0.48	1.00	ND	Pass	R1 M1	Butoxide	0.70	2.00	LUQ	F d55	
Daminozide	0.48	1.00	ND	Pass	L1 M1	Prallethrin	0.10	0.20	ND	Pass	R1 M1
Diazinon	0.10	0.20	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Pyrethrins	0.48	1.00	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass		Spinosad	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
Flonicamid	0.48	1.00	ND	Pass		Thiacloprid	0.10	0.20	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass		Thiamethoxam	0.10	0.20	ND	Pass	
						Trifloxystrobin	0.10	0.20	ND	Pass	

Instrument: LC-QQQ ; Method: TPL_Pesticides_01



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Pass

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Mac 1 (Caps Cut) Flower/Pre-Roll

Plant, Flower - Cured

 $\label{eq:listense} Dispensary\, License\, \#: ;\, Cultivation\, License\, H: ;\, Cultivati$

Heavy Metals

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	L1 V1
Cadmium	200.0	400.0	<loq< th=""><th>Pass</th><th>L1 V1</th></loq<>	Pass	L1 V1
Lead	500.0	1000.0	ND	Pass	L1 V1
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>L1V1</th></loq<>	Pass	L1V1

Microbials				Pass
Analyte	LOQ	Limit	Result	StatusQualifier
	CFU/g	CFU/g	CFU/g	
E. Coli	10	100	<10	Pass

Analyte	Limit	Result	Status	Qualifier
Salmonella	Detectable in 1g	Not Detected	Pass	
Aspergillus	Detectable in 1g	Not Detected	Pass	
Aspergillus fumigatus	Detectable in 1g	Not Detected	Pass	
Aspergillus niger	Detectable in 1g	Not Detected	Pass	
Aspergillus flavus	Detectable in 1g	Not Detected	Pass	
Aspergillus terreus	Detectable in 1g	Not Detected	Pass	

Instrument: ICPMS; Method: AOAC 2021.03

Instrument: qPCR/Plating; AOAC Methods 082102, 022202 and 2018.13



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B1 = Target analyte detected in calibration blank was above LOQ but the concentration of cannabinoid was blow LOQ,

B2 = Target analyte detected in calibration blank was above LOQ but was below the maximum allowable concentration.

D1 = The limit of quantitation and the sample results were adjusted to reflect sample dilution,

11 = The relative intensity of a characteristic ion in a sample analyte exceeded the acceptance criteria with respect to the reference spectra, indicating interference,

L1 = The percent recovery of a laboratory control sample is greater than the acceptance limits in A.A.C 17 R9-17-404.03(K)(2)(C), but the sample's target analytes were not detected above the maximum allowed concentration,

M1 = The recovery from the matrix spike was high, but the recovery from the laboratory control sample was within acceptance criteria,

M2 = The recovery from the matrix spike was low, but the recovery from the laboratory control sample was within acceptance criteria,

M3 = The recovery from the matrix spike was unusable because the analyte concentration was disproportionate to the spike level, but the recovery from the laboratory control sample was within acceptance criteria,

M4 = The analysis of a spiked sample required a dilution such that the spike recovery calculation does not provide useful information, but the recovery from the associated laboratory control sample was within acceptance criteria,

M5 = The analyte concentration was determined by the method of standard addition, in which the standard is added directly to the aliquots of the analyzed sample,

N1 - A description of the variance is described in the final report of testing,

R1 = The relative percent difference for the laboratory control sample and duplicate exceeded the limit in A.A.C 17 R9-17-404.03(K)(3), but the recover in subsection A.A.C 17 R9-17-404.03 (K)(2) was within accepted criteria,

R2 = The relative percent difference for a sample and duplicated exceeded the limit in subsection A.A.C 17 R9-17-404.03 (O)

Q1 = Sample integrity was not maintained,

Q2 = The sample is heterogenous and sample homogeneity could not be readily achieved using routine laboratory practices

Q3 = Testing result is for informational purposes only and cannot be used to satisfy dispensary testing requirements in R9-17-317.01(A) or labeling requirements in R9-17-317

V1 = The recovery from continuing calibration verification standards exceeded the acceptance limits denoted in A.C.C 17 R9-17-403.03(J)(1)(b), but the sample's target analytes were not detected above the maximum allowable concentrations for the analytes in the sample.

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