Certificate of Analysis

Genesis Bioceuticals, LLC

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Phoenix, AZ 85007
shonae.j@genbioaz.com
(847) 682-4899
Lic. #00000058DCQU00115543
Harvest Dates:

Grape Illusion Shatter

Concentrates & Extracts, Shatter, Extraction Method: Butane
Dispensary License #: ; Manufacturing License #: ; Cultivation License #:

Sample: 2312TLL0063.0165

Strain: Grape Illusion
Parent Batch #: ; Batch#: GILL-1871-20230516-S; Batch Size: 11 g
Sample Received: 12/11/2023; Report Created: 12/13/2023; Expires: 12/13/2024
Manufacturing Date:
Sampling: ; Environment:

Safety

Pass Pesticides	**Pass** Microbials	**Pass** Mycotoxins
Pass Solvents	**Pass** Metals	**Not Tested** Foreign Matter

Cannabinoids

TPL_Potency_01

85.75% Total THC	0.16% Total CBD	NT Moisture

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	95.95	959.5	
Δ9-THC	0.10	1.60	16.0	
Δ8-THC	0.10	NR	NR	
THCV	0.10	NR	NR	
CBDa	0.10	0.18	1.8	
CBD	0.10	NR	NR	
CBDV	0.10	NR	NR	
CBN	0.10	NR	NR	
CBGa	0.10	0.93	9.3	
CBG	0.10	0.34	3.4	
CBC	0.10	NR	NR	
Total		**99.01**	**990.1**	

Total THC = THCa * 0.877 + Δ9-THC
Total CBD = CBDa * 0.877 + CBD
Instrument: HPLC-DAD: ; Method: TPL_Potency_01

Terpenes

TPL_Terpenes_01

Hops	Cinnamon	Earthy

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
α-Humulene		2.3020	23.020	
β-Caryophyllene		1.9870	19.870	
Ocimene		0.7580	7.580	
Linalool		0.5600	5.600	
trans-Nerolidol		0.5550	5.550	
β-Pinene		0.3820	3.820	
α-Bisabolol		0.2760	2.760	
β-Myrcene		0.2170	2.170	
α-Pinene		0.1760	1.760	
Terpinolene		0.1370	1.370	
γ-Terpinene		0.1080	1.080	
Camphene		0.0600	0.600	
Isopulegol		0.0150	0.150	
3-Carene		<	<	
α-Terpinene		<	<	
cis-Nerolidol		<	<	
Geraniol		<	<	
Guaiol		<	<	
Total		**7.5330**	**75.330**	

Instrument: GCMS ; Method: TPL_Terp_01
Notes:

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Brian DiMarco
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Residual Solvents — Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Acetone	196.0	1000.0	ND	Pass	
Acetonitrile	80.0	410.0	ND	Pass	
Benzene	0.4	2.0	ND	Pass	
n-Butane			NR	NT	
Isobutane			NR	NT	
Butanes	490.0	5000.0	<LOQ	Pass	
Chloroform	12.0	60.0	ND	Pass	
Dichloromethane	118.0	600.0	ND	Pass	
Ethanol	980.0	5000.0	ND	Pass	
Ethyl-Acetate	980.0	5000.0	ND	Pass	
Ethyl-Ether	980.0	5000.0	ND	Pass	
Heptane	980.0	5000.0	ND	Pass	
n-Hexane			NR	NT	
2-Methyl-Pentane			NR	NT	
3-Methyl-Pentane			NR	NT	
2,2-Dimethyl-Butane			NR	NT	
2,3-Dimethyl-Butane			NR	NT	
Hexanes	142.0	290.0	ND	Pass	
Isopropyl-Acetate	980.0	5000.0	ND	Pass	
Methanol	588.0	3000.0	ND	Pass	
n-Pentane			NR	NT	
Isopentane			NR	NT	
Neopentane			NR	NT	
Pentanes	980.0	5000.0	ND	Pass	
2-Propanol	980.0	5000.0	<LOQ	Pass	
Propane			NR	NT	
Toluene	175.0	890.0	ND	Pass	
1,2-Dimethyl-Benzene			NR	NT	
1,3-Dimethyl-Benzene			NR	NT	
1,4-Dimethyl-Benzene			NR	NT	
Ethyl-Benzene			NR	NT	
Xylenes	851.0	2170.0	ND	Pass	

Instrument: HS-GCMS

Heavy Metals — Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	
Cadmium	200.0	400.0	ND	Pass	
Lead	500.0	1000.0	ND	Pass	
Mercury	100.0	200.0	<LOQ	Pass	M2

Mycotoxins — Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
B1	10	0	ND	Pass	
B2	10	0	ND	Pass	
G1	10	0	ND	Pass	
G2	5	0	ND	Pass	I1,V1
Ochratoxin A	10	20	ND	Pass	I1,M1
Total Aflatoxins	9	20	ND	Pass	V1

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Microbials

Pass

Analyte	LOQ	Limit	Result	Status	Qualifier
	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: Plating/qPRC ; Method: AOAC 82102 and 022202

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Grape Illusion Shatter

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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	
Acephate	0.19	0.40	ND	Pass	
Acequinocyl			NR	NT	
Acetamiprid	0.09	0.20	ND	Pass	
Aldicarb	0.19	0.40	ND	Pass	
Azoxystrobin	0.09	0.20	ND	Pass	
Bifenazate	0.09	0.20	ND	Pass	L1
Bifenthrin	0.09	0.20	ND	Pass	L1,V1
Boscalid	0.19	0.40	ND	Pass	M2
Carbaryl	0.09	0.20	ND	Pass	
Carbofuran	0.09	0.20	ND	Pass	
Chlorantraniliprole	0.09	0.20	ND	Pass	L1
Chlorfenapyr	0.47	1.00	ND	Pass	I1
Chlorpyrifos	0.09	0.20	ND	Pass	
Clofentezine	0.09	0.20	ND	Pass	M2
Cyfluthrin	0.47	1.00	ND	Pass	M2
Cypermethrin	0.47	1.00	ND	Pass	M2,V1
Daminozide	0.47	1.00	ND	Pass	
Diazinon	0.09	0.20	ND	Pass	
Dichlorvos	0.02	0.10	ND	Pass	
Dimethoate	0.09	0.20	ND	Pass	
Ethoprophos	0.09	0.20	ND	Pass	L1
Etofenprox	0.19	0.40	ND	Pass	L1,V1
Etoxazole	0.09	0.20	ND	Pass	
Fenoxycarb	0.09	0.20	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass	M2
Fipronil	0.19	0.40	ND	Pass	I1,R1
Flonicamid	0.47	1.00	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass	M2

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Hexythiazox	0.47	1.00	ND	Pass	M2
Imazalil	0.09	0.20	ND	Pass	I1
Imidacloprid	0.19	0.40	ND	Pass	
Kresoxim Methyl	0.19	0.40	ND	Pass	M2
Malathion	0.09	0.20	ND	Pass	
Metalaxyl	0.09	0.20	ND	Pass	
Methiocarb	0.09	0.20	ND	Pass	L1
Methomyl	0.19	0.40	ND	Pass	
Myclobutanil	0.09	0.20	ND	Pass	
Naled	0.24	0.50	ND	Pass	
Oxamyl	0.47	1.00	ND	Pass	
Paclobutrazol	0.19	0.40	ND	Pass	L1,V1
Permethrin	0.09	0.20	ND	Pass	M2,V1
Phosmet	0.09	0.20	ND	Pass	
Piperonyl Butoxide	0.94	2.00	ND	Pass	M2
Prallethrin	0.09	0.20	ND	Pass	
Propiconazole	0.19	0.40	ND	Pass	L1,V1
Propoxur	0.09	0.20	ND	Pass	
Pyrethrins	0.47	1.00	ND	Pass	
Pyridaben	0.09	0.20	ND	Pass	M2
Spinosad	0.09	0.20	ND	Pass	M2
Spiromesifen	0.09	0.20	ND	Pass	
Spirotetramat	0.09	0.20	ND	Pass	
Spiroxamine	0.19	0.40	ND	Pass	
Tebuconazole	0.19	0.40	ND	Pass	L1,V1
Thiacloprid	0.09	0.20	ND	Pass	
Thiamethoxam	0.09	0.20	ND	Pass	
Trifloxystrobin	0.09	0.20	ND	Pass	M2

Instrument: LC-QQQ

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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,

I1 = 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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