

## Genesis Biocenticals, LLC

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(847) 682-4899  
Lic. #00000058DCQU00115543  
Harvest Dates: 12/06/2023

## Sample: 2402TLL0051.0299

Strain: Strawberries n Cream  
Parent Batch #: ; Batch#: G-0207-SC; Batch Size: 16 g  
Sample Received: 02/13/2024; Report Created: 02/19/2024; Expires: 02/19/2025  
Manufacturing Date: 02/07/2024  
Sampling: ; Environment:

## Strawberries n Cream Cured Resin Batter

Concentrates & Extracts, Batter/Badder, Extraction Method: Butane  
Dispensary License #: ; Manufacturing License #: ; Cultivation License #:



## Safety

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

## Cannabinoids

TPL\_Potency\_01

83.42%	0.10%	99.42%
Total THC	Total CBD	Total Cannabinoids Q3

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	93.83	938.3	
Δ9-THC	0.10	1.13	11.3	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	0.12	1.2	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	4.17	41.7	
CBG	0.10	0.17	1.7	
CBC	0.10	ND	ND	
<b>Total</b>		<b>99.42</b>	<b>994.2</b>	

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
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Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
α-Humulene		1.6300	16.300	Q3
β-Caryophyllene		1.3410	13.410	Q3
Ocimene		0.6810	6.810	Q3
δ-Limonene		0.6330	6.330	Q3
trans-Nerolidol		0.3860	3.860	Q3
β-Pinene		0.2570	2.570	Q3
β-Myrcene		0.1900	1.900	Q3
Terpinolene		0.1530	1.530	Q3
α-Pinene		0.1470	1.470	Q3
γ-Terpinene		0.1330	1.330	Q3
Linalool		0.1230	1.230	Q3
α-Bisabolol		0.1160	1.160	Q3
Eucalyptol		0.1050	1.050	Q3
Caryophyllene Oxide		0.0990	0.990	Q3
cis-Nerolidol		0.0860	0.860	Q3
Camphene		0.0270	0.270	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
<b>Total</b>		<b>6.1070</b>	<b>61.070</b>	

Instrument: GCMS; Method: TPL\_Terp\_01  
Notes:

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## Pesticides TPL\_Pesticides\_01

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier	Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	R1 M1	Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass	R1	Imazalil	0.10	0.20	ND	Pass	R1 L1
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	R1
Aldicarb	0.19	0.40	ND	Pass	R1 L1	Kresoxim	0.19	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass	R1 L1	Methyl					
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	R1
Bifenthrin	0.10	0.20	ND	Pass	R1	Metalaxyl	0.10	0.20	ND	Pass	R1 L1
Boscalid	0.19	0.40	ND	Pass	R1	Methiocarb	0.10	0.20	ND	Pass	R1 L1
Carbaryl	0.10	0.20	ND	Pass	R1	Methomyl	0.19	0.40	ND	Pass	L1
Carbofuran	0.10	0.20	ND	Pass	R1	Myclobutanil	0.10	0.20	ND	Pass	R1
Chlorantraniliprole	0.10	0.20	ND	Pass	R1	Naled	0.24	0.50	ND	Pass	R1
Chlorfenapyr	0.48	1.00	ND	Pass	R1 M2	Oxamyl	0.48	1.00	ND	Pass	R1
Chlorpyrifos	0.10	0.20	ND	Pass	R1	Paclobutrazol	0.19	0.40	ND	Pass	L1
Clofentezine	0.10	0.20	ND	Pass	R1 L1	Permethrin	0.10	0.20	ND	Pass	R1
Cyfluthrin	0.48	1.00	ND	Pass	R1 L1	Phosmet	0.10	0.20	ND	Pass	R1
Cypermethrin	0.48	1.00	ND	Pass	R1 L1	Piperonyl Butoxide	0.96	2.00	ND	Pass	R1 L1
Daminozide	0.48	1.00	ND	Pass	R1 M2	Prallethrin	0.10	0.20	ND	Pass	R1 L1
Diazinon	0.10	0.20	ND	Pass	L1	Propiconazole	0.19	0.40	ND	Pass	R1
Dichlorvos	0.05	0.10	ND	Pass	R1	Propoxur	0.10	0.20	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass	R1	Pyrethrins	0.48	1.00	ND	Pass	R1 L1
Ethoprophos	0.10	0.20	ND	Pass	R1 L1	Pyridaben	0.10	0.20	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass	R1 L1	Spinosad	0.10	0.20	ND	Pass	
Etoazole	0.10	0.20	ND	Pass	R1	Spiromesifen	0.10	0.20	ND	Pass	R1 L1
Fenoxycarb	0.10	0.20	ND	Pass	R1 L1	Spirotetramat	0.10	0.20	ND	Pass	R1 L1
Fenpyroximate	0.19	0.40	ND	Pass	R1 L1	Spiroxamine	0.19	0.40	ND	Pass	R1 L1
Fipronil	0.19	0.40	ND	Pass	R1	Tebuconazole	0.19	0.40	ND	Pass	R1 L1
Fonicamid	0.48	1.00	ND	Pass	R1 L1	Thiacloprid	0.10	0.20	ND	Pass	R1 L1
Fludioxonil	0.19	0.40	ND	Pass	R1	Thiamethoxam	0.10	0.20	ND	Pass	R1
						Trifloxystrobin	0.10	0.20	ND	Pass	

Instrument: LC-QQQ ; Method: TPL\_Pesticides\_01

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### Heavy Metals Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	
Cadmium	200.0	400.0	<LOQ	Pass	M1
Lead	500.0	1000.0	<LOQ	Pass	M1
Mercury	100.0	200.0	<LOQ	Pass	

### Microbials Pass

Analyte	LOQ	Limit	Result	Status	Qualifier
	CFU/g	CFU/g	CFU/g		
E. Coli	10	100	<10	Pass	

### Residual Solvents Pass

Instrument: ICPMS; Method: AOAC 2021.03

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Acetone	199.0	1000.0	ND	Pass	
Acetonitrile	82.0	410.0	ND	Pass	
Benzene	0.4	2.0	ND	Pass	
Butanes	498.0	5000.0	ND	Pass	
Chloroform	12.0	60.0	ND	Pass	
Dichloromethane	120.0	600.0	ND	Pass	
Ethanol	996.0	5000.0	ND	Pass	
Ethyl-Acetate	996.0	5000.0	ND	Pass	
Ethyl-Ether	996.0	5000.0	ND	Pass	
Heptane	996.0	5000.0	ND	Pass	
Hexanes	144.0	290.0	ND	Pass	
Isopropyl-Acetate	996.0	5000.0	ND	Pass	
Methanol	598.0	3000.0	ND	Pass	
Pentanes	996.0	5000.0	ND	Pass	
2-Propanol	996.0	5000.0	ND	Pass	
Toluene	177.0	890.0	ND	Pass	
Xylenes	865.0	2170.0	ND	Pass	

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

### Mycotoxins Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
B1	8	20	ND	Pass	
B2	8	20	ND	Pass	
G1	8	20	ND	Pass	
G2	8	20	ND	Pass	
Ochratoxin A	8	20	ND	Pass	
Total Aflatoxins	8	20	ND	Pass	

Instrument: HS-GCMS; Method: TPL\_ResSolv\_01

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