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RSP Productions: Asylum

2985 W. Whitton Ave. Phoenix, AZ 85017 Iodellrichardson@gmail.com (702) 338-7088 Lic. #00000070DCBD00783295 Sample: 2405TLL0154.0769

Strain: Grape Gas

Parent Batch #:; Batch#: 20240423R8GG; Batch Size: 15 g

Sample Received: 05/03/2024; Report Created: 05/10/2024; Expires: 05/10/2025

Manufacturing Date:

Sampling: ; Environment:

Harvest Dates: 04/23/2024 Grape Gas Flower

Plant, Flower - Cured

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





Safety

Pass

Pesticides

Pass

Microbials

Pass

Metals

Cannabinoids

TPL_Potency_01

26.40% Total THC ND Total CBD

31.23%Total Cannabinoids
Q3

Analyte	LOQ	Mass	Mass	Qualifie
	%	%	mg/g	
THCa	0.10	29.57	295.7	
Δ9-THC	0.10	0.47	4.7	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	ND	ND	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	1.20	12.0	
CBG	0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
CBC	0.10	ND	ND	
Total		31.23	312.3	

Terpenes TPL_Terpenes_01



Lemon

Earthy



Mass Qualifier Analyte Mass 0.2400 mg/g 2.400 δ-Limonene 0.2300 2.300 Ocimene β-Caryophyllene α-Humulene 0.2000 2.000 0.1600 1.600 ß-Myrcene 0.1200 1.200 β-Pinene 0.1200 1.200 0.1000 1.000 Linalool 0.0800 0.800 Terpinolene 0.0700 0.700 y-Terpinene trans-Nerolidol 0.0700 0.700 0.0600 0.600 α-Pinene α-Bisabolol 0.0200 0.200 0.0100 0.100 Camphene 3-Carene α-Terpinene Caryophyllene Oxide cis-Nerolidol Eucalyptol Geraniol Guaiol Isopulegol p-Cymene Total 1.4800 14.800

Instrument: GCMS; Method: TPL_Terp_01

Notes

Total THC = THCa * 0.877 + Δ 9-THC Total CBD = CBDa * 0.877 + CBD

Instrument: HPLC-DAD: ; Method: TPL_Potency_01



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Grape Gas Flower

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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status C	(ualifier	Analyte	LOQ	Limit	Mass	Status (Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1L1	Hexythiazox	0.48	1.00	ND	Pass	L1 V1
Abameetin					V11	lmazalil	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	
Aldicarb	0.19	0.40	ND	Pass		Methyl					
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenazate	0.10	0.20	ND	Pass	M1V1	Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	L1 V1	Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass	V1	Methomyl	0.19	0.40	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	V1
Carbofuran	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass	7.0	Oxamyl	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	M2	Paclobutrazol	0.19	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	L1 V1
Clofentezine	0.10	0.20	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass	M1L1	Piperonyl	0.96	2.00	<loq< td=""><td>Pass</td><td></td></loq<>	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1L1	Butoxide	0.40	0.00		_	14114
Daminozide	0.48	1.00	ND		R1 M1	Prallethrin	0.10 0.19	0.20	ND ND	Pass Pass	M1V1
Diazinon	0.40	0.20	ND	Pass Pass	IVII	Propiconazole Propoxur	0.19	0.40	ND	Pass	
Dichloryos	0.10	0.20	ND	Pass		Pyrethrins	0.10	1.00	ND	Pass	V1
Dimethoate	0.03	0.10	ND	Pass		Pyridaben	0.40	0.20	ND	Pass	V1 V1
Ethoprophos	0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	V 1
Etofenprox	0.10	0.20	ND	Pass	V1	Spiromesifen	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.40	ND	Pass	A T	Spirotetramat	0.10	0.20	ND	Pass	М1
Fenoxycarb	0.10	0.20	ND	Pass	M2 V1	Spiroxamine	0.10	0.20	ND	Pass	1417
Fenpyroximate	0.10	0.40	ND	Pass	L1	Tebuconazole	0.17	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass	-1	Thiacloprid	0.10	0.20	ND	Pass	
Flonicamid	0.48	1.00	ND	Pass	V1	Thiamethoxam	0.10	0.20	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass	• •	Trifloxystrobin	0.10	0.20	ND	Pass	

 $Instrument: LC\text{-}QQQ \ ; Method: TPL_Pesticides_01$



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Heavy Me	tals				Pass
Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	V1
Cadmium	200.0	400.0	<loq< th=""><th>Pass</th><th></th></loq<>	Pass	
Lead	500.0	1000.0	<loq< th=""><th>Pass</th><th></th></loq<>	Pass	
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>L1 R1</th></loq<>	Pass	L1 R1

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	<u> </u>
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	

LOQ=Limit of Quantitation. The reported result is based on a simple weight with the applicable moisture content for that sample. Unless otherwise stated, all quality control samples performed within specifications established by the Laboratory. 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,
- 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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