

## RSP Productions: Asylum

2985 W. Whitton Ave.  
Phoenix, AZ 85017  
lodellrichardson@gmail.com  
(702) 338-7088  
Lic. #00000070DCBD00783295  
Harvest Dates: 10/01/2024

## Sample: 2410TLL0352.1628

Strain: GMO  
Parent Batch #: ; Batch#: 20241001R4GMO; Batch Size: g  
Sample Received: 10/04/2024; Report Created: 10/10/2024  
Manufacturing Date:  
Sampling: ; Environment:

## GMO Flower

Plant, Flower - Cured, Extraction Method: Indoor  
Dispensary License #: ; Manufacturing License #: ; Cultivation License #:



## Safety

<b>Pass</b>	<b>Pass</b>	<b>Pass</b>
Pesticides	Microbials	Metals

## Cannabinoids

TPL\_Potency\_01

<b>30.09%</b>	<b>&lt;LOQ</b>	<b>35.21%</b>
Total THC	Total CBD	Total Cannabinoids Q3

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	32.90	329.0	
Δ9-THC	0.10	1.24	12.4	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	<LOQ	<LOQ	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	0.87	8.7	
CBG	0.10	0.20	2.0	
CBC	0.10	ND	ND	
<b>Total</b>		<b>35.21</b>	<b>352.1</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

Mint	Wood	Turpentine

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
Isopulegol		1.0750	10.750	Q3
Caryophyllene Oxide		1.0550	10.550	Q3
Terpinolene		0.3170	3.170	Q3
γ-Terpinene		0.1840	1.840	Q3
α-Terpinene		0.1650	1.650	Q3
Geraniol		0.1580	1.580	Q3
Camphene		0.0660	0.660	Q3
α-Pinene		0.0460	0.460	Q3
β-Caryophyllene		0.0270	0.270	Q3
cis-Nerolidol		0.0090	0.090	Q3
3-Carene		<	<	Q3
α-Bisabolol		<	<	Q3
α-Humulene		<	<	Q3
β-Myrcene		<	<	Q3
β-Pinene		<	<	Q3
δ-Limonene		<	<	Q3
Eucalyptol		<	<	Q3
Guaiol		<	<	Q3
Linalool		<	<	Q3
Ocimene		<	<	Q3
p-Cymene		<	<	Q3
trans-Nerolidol		<	<	Q3
<b>Total</b>		<b>3.1020</b>	<b>31.020</b>	

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

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## Pesticides TPL\_Pesticides\_01

Pass

Analyte	LOQ	Limit	Result	Status	Qualifier	Analyte	LOQ	Limit	Result	Status	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass		Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass		Imazalil	0.10	0.20	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
Aldicarb	0.19	0.40	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass		Methyl					
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass		Metalaxyl	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Methomyl	0.19	0.40	ND	Pass	
Carbofuran	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass		Oxamyl	0.48	1.00	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Pacllobutrazol	0.19	0.40	ND	Pass	
Clofentezine	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass		Piperonyl					
Daminozide	0.48	1.00	ND	Pass		Butoxide	0.96	2.00	ND	Pass	
Diazinon	0.10	0.20	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Pyrethrins	0.48	1.00	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
Fipronil	0.19	0.40	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	
Fonicamid	0.48	1.00	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass		Thiacloprid	0.10	0.20	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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### 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	L1
Lead	500.0	1000.0	ND	Pass	L1
Mercury	100.0	200.0	<LOQ	Pass	L1

### 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	

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

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

Brian DiMarco  
Laboratory Director

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coa.support@confidentlims.com  
(866) 506-5866  
www.confidentlims.com



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 recovery 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 heterogeneous 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.A.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.