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Sample: 2408TLL0276.1309

### **RSP** Productions: Asylum

2985 W. Whitton Ave. Phoenix, AZ 85017 Iodellrichardson@gmail.com (702) 338-7088 Lic. #00000070DCBD00783295 Harvest Dates: 07/30/2024 Isis x ONYCD Flower Strain: Isis x ONYCD Parent Batch #: ; Batch#: 20240730R4ISO; Batch Size: g Sample Received: 08/02/2024; Report Created: 08/07/2024; Expires: 08/07/2025 Manufacturing Date: Sampling: ; Environment:







#### Cannabinoids

TPL_Potency_01		_		
20.50%	<loq< th=""><th></th><th>23.79</th><th>%</th></loq<>		23.79	%
Total THC	Total CBD		Total Canna Q3	binoids
Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	22.53	225.3	
Δ9-THC	0.10	0.75	7.5	
∆8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	0.41	4.1	
CBG	0.10	0.11	1.1	
CBC	0.10	ND	ND	
Total		23.79	237.9	

Total THC = THCa \*  $0.877 + \Delta 9$ -THC Total CBD = CBDa \* 0.877 + CBDInstrument: HPLC-DAD: ; Method: TPL\_Potency\_01



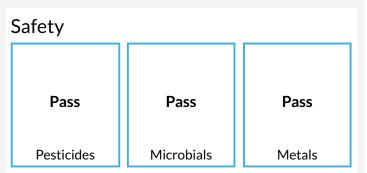
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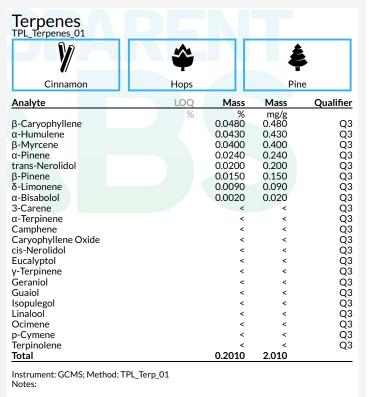
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Brian DiMarco Laboratory Director Confident LIMS All Rights Reserved coa.support@confidentlims.com (866) 506-5866 www.confidentlims.com



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#### **RSP** Productions: Asylum

2985 W. Whitton Ave. Phoenix, AZ 85017 Iodellrichardson@gmail.com Sample (702) 338-7088 Lic. #0000070DCBD00783295 Harvest Dates: 07/30/2024 Isis x ONYCD Flower

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### Pesticides TPL\_Pesticides\_01

### Sample: 2408TLL0276.1309

Strain: Isis x ONYCD Parent Batch #: ; Batch#: 20240730R4ISO; Batch Size: g Sample Received: 08/02/2024; Report Created: 08/07/2024; Expires: 08/07/2025 Manufacturing Date: Sampling: ; Environment:



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	0.19	0.40	ND	Pass
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	Paclobutrazol	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	0.96	2.00	ND	Pass
Daminozide	0.48	1.00	ND	Pass	Butoxide	0.70	2.00	ND	Fass
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	0.77	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
Flonicamid	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
									_

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



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Trifloxystrobin

0.10

0.20

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ND

Pass



Brian DiMarco Laboratory Director

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Pass

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Strain: Isis x ONYCD

Sample: 2408TLL0276.1309

Parent Batch #:; Batch#: 20240730R4ISO; Batch Size: g

#### **RSP** Productions: Asylum

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**Heavy Metals** 

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	
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></th></loq<>	Pass	

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	

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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Laboratory Director

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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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