

## Certificate of Analysis



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

2985 W. Whitton Ave. Phoenix, AZ 85017 lodellrichardson@gmail.com (702) 338-7088 Lic. #00000070DCBD00783295 Harvest Dates: 09/24/2024

### Doc Holiday Flower

Plant, Flower - Cured, Extraction Method: Indoor Dispensary License #:; Manufacturing License #:; Cultivation License #: Sample: 2409TLL0343.1607

Strain: Doc Holiday

Parent Batch #:; Batch#: 20240924R3DH; Batch Size: g Sample Received: 09/30/2024; Report Created: 10/02/2024

Manufacturing Date:

Sampling: ; Environment:





#### Safety

**Pass Pesticides** 

**Pass** Microbials

**Pass** 

Metals

#### Cannabinoids

TPL\_Potency\_01

25.33% Total THC

<LOQ Total CBD To

29.59%	l
otal Cannabinoids Q3	ı

II L_ICIP	ches_or	
	Mint	
Analuta		





Analyte	LOQ	Mass	Mass	Qualifie
	%	%	mg/g	
THCa	0.10	27.32	273.2	
Δ9-THC	0.10	1.36	13.6	
Δ8-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	0.76	7.6	
CBG	0.10	0.14	1.4	
CBC	0.10	ND	ND	
Total		29.59	295.9	

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

LABS

Terpenes





Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
Isopulegol		0.5090	5.090	Q3
Caryophyllene Oxid	e	0.5020	5.020	Q3
y-Terpinene		0.3160	3.160	Q3
Terpinolene		0.2150	2.150	Q3
Geraniol		0.1330	1.330	Q3
α-Terpinene		0.1120	1.120	Q3
α-Bisabolol		0.0930	0.930	Q3
α-Pinene		0.0440	0.440	Q3
Camphene		0.0420	0.420	Q3
β-Caryophyllene		0.0260	0.260	Q3
cis-Nerolidol		0.0050	0.050	Q3
3-Carene		<	<	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
Total		1.9970	19.970	

Instrument: GCMS; Method: TPL\_Terp\_01

1721 E McDowell Road Phoenix, AZ

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Brian DiMarco **Laboratory Director** 

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#### Pesticides TPL\_Pesticides\_01

Analyte	LOQ	Limit	Result	Status Q	ualifier	Analyte	LOQ	Limit	Result	Status Q	ualifier
	PPM	PPM	PPM		<u></u>		PPM	PPM	PPM		<u>.</u>
Abamectin	0.24	0.50	ND	Pass	V1	Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass		lmazalil	0.10	0.20	ND	Pass	R1
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	V1
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	F 455	
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	V1
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	V1L1	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	V1
						Trifloxystrobin	0.10	0.20	ND	Pass	

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



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Lead

Mercury

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3 of 4

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Heavy Me	tals				Pass
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>L1</th></loq<>	Pass	L1

<LOQ

<LOQ

Pass

Pass

L1

L1

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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- 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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