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Sample: 2401TLL0036.0230

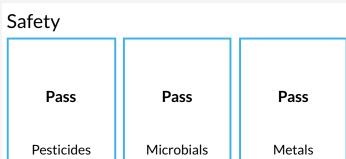
## Total Health & Wellness dba True Harvest

Phoenix, AZ 85043 jpastor@trueharvestco.com Strain: Anslinger's Demise Parent Batch #: ; Batch#: R4AND0109; Batch Size: 20 g Sample Received: 01/31/2024; Report Created: 02/02/2024; Expires: 02/02/2025 Manufacturing Date: Sampling: ; Environment:

Lic. #00000100DCWU00857159 Harvest Dates:

### Anslinger's Demise

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



Hops

Mass

0.6190

0.6080

0.6010

0.5380

0.5130

0.2300

0.2070

0.1650

0.1000

0.0970

0.0750

100



### Cannabinoids

	_		
ND		32.99%	
Total CBD		Total Cannabinoid Q3	
LOQ	Mass	Mass	Qualifier
%	%	mg/g	
0.10	31.89	318.9	
0.10	0.14	1.4	
0.10	ND	ND	
0.10	0.96	9.6	
0.10	ND	ND	
0.10	ND	ND	
	32.99	329.9	
	Total CBD   %   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10   0.10	Total CBD   LOQ Mass   % %   0.10 31.89   0.10 0.14   0.10 ND   0.10 ND	Total CBD Total Canna Q3   LOQ Mass Mass   % % mg/g   0.10 31.89 318.9   0.10 0.14 1.4   0.10 ND ND   0.10 ND ND

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



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0.0670 y-Terpinene α-Bisabolol 0.0550 Eucalyptol 0.0460 0.0230 Camphene Caryophyllene Oxide 0.0120 α-Terpinene cis-Nerolidol Geraniol Guaiol Isopulegol p-Cymene Total 3.9560

Instrument: GCMS; Method: TPL\_Terp\_01

Notes:

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(866) 506-5866



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Lavender

Mass mg/g 6.190

6.080

6.010

5.380

5.130

2.300

2.070

1.650

1.000

0.970

0.750

0.670

0.550

0.460

0.230

0.120

39.560

<

Qualifier

Q3

Q3

Terpenes TPL\_Terpenes\_01

Lemon

Analyte

Linalool Ocimene

δ-Limonene

α-Humulene

β-Myrcene

. Terpinolene

ß-Pinene

3-Carene

α-Pinene

β-Caryophyllene trans-Nerolidol



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### Pesticides TPL Pesticides 01

Analyte	LOQ	Limit	Mass	Status C	Qualifier	Analyte	LOQ	Limit	Mass	Status C	ualifier
· · ·	PPM	PPM	PPM		•		PPM	PPM	PPM		·
Abamectin	0.24	0.50	ND	Pass M	11   1 V 1	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	M1	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	M2
Cyfluthrin	0.48	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1	Piperonyl	0.96	2.00	ND	Pass	
Daminozide	0.48	1.00	ND	Pass	M1L1	Butoxide	0.70	2.00	ND	rd55	
Daminozide	0.40	1.00	ND	Pass	V1	Prallethrin	0.10	0.20	ND	Pass	
Diazinon	0.10	0.20	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Pyrethrins	0.48	1.00	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass		Spinosad	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
Flonicamid	0.48	1.00	ND	Pass		Thiacloprid	0.10	0.20	ND	Pass	
Fludioxonil	0.19	0.40	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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**Brian DiMarco** 

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Pass

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### **Heavy Metals**

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></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	

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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