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True Harvest Sample: 2401TLL0006.0064

Strain: Lemon Shiv

Phoenix, AZ 85043 Parent Batch #:; Batch#: R7LS1128; Batch Size: 20 undefined mike@trueharvestco.com Sample Received: 01/10/2024; Report Created: 01/15/2024; Expires: 01/15/2025

Manufacturing Date:

Sampling: ; Environment:

Harvest Dates: 2023-11-28 Lemon Shiv

Lic. #00000100DCWU00857159

Plant, Flower - Cured

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





Safety

Pass **Pesticides**

Microbials **Mycotoxins**

Not Tested Solvents

Pass Metals

Pass

Not Tested

Not Tested

Foreign Matter

Cannabinoids

TPL_Potency_01

25.26% **Total THC**

<LOQ Total CBD

30.16% **Total Cannabinoids**

LOQ	Mass	Mass	Qualifie
%	%	mg/g	
0.20	27.62	276.2	
0.20	1.03	10.3	
0.20	ND	ND	
0.20	ND	ND	
0.20	<loq< th=""><th><loq< th=""><th></th></loq<></th></loq<>	<loq< th=""><th></th></loq<>	
0.20	ND	ND	
0.20	ND	ND	
0.20	ND	ND	
0.20	1.51	15.1	
0.20	<loq< th=""><th><loq< th=""><th></th></loq<></th></loq<>	<loq< th=""><th></th></loq<>	
0.20	ND	ND	
	30.16	301.6	
	% 0.20 0.20 0.20 0.20 0.20 0.20 0.20 0.2	% % 0.20 27.62 0.20 1.03 0.20 ND	% % mg/g 0.20 27.62 276.2 0.20 1.03 10.3 0.20 ND ND 0.20 ND ND 0.20 <loq< td=""> <loq< td=""> 0.20 ND ND 0.20 ND ND 0.20 ND ND 0.20 ND ND 0.20 1.51 15.1 0.20 <loq< td=""> <loq< td=""> 0.20 ND ND</loq<></loq<></loq<></loq<>

Terpenes TPL_Terpenes_01









Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
Terpinolene		0.629	6.29	Q3
trans-Nerolidol		0.311	3.11	Q3
Ocimene		0.277	2.77	Q3
α-Humulene		0.221	2.21	Q3
β-Caryophyllene		0.221	2.21	Q3
δ-Limonene		0.196	1.96	Q3
β-Pinene		0.161	1.61	Q3
β-Myrcene		0.152	1.52	Q3
3-Carene		0.131	1.31	Q3
α-Pinene		0.094	0.94	Q3
α-Bisabolol		0.090	0.90	Q3
Guaiol		0.084	0.84	Q3
y-Terpinene		0.065	0.65	Q3
α-Terpinene		0.061	0.61	Q3
Linalool		0.050	0.50	Q3
Eucalyptol		0.045	0.45	Q3
Camphene		0.011	0.11	Q3
Caryophyllene Ox	ide	0.010	0.10	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		2.809	28.09	

Instrument: GCMS; Method: TPL_Terp_01

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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Brian DiMarco Laboratory Director

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Harvest Dates: 2023-11-28 **Lemon Shiv**

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Microbials				Pass
Analyte	LOQ	Limit	Result	StatusQualifier
	CFU/g	CFU/g	CFU/g	
E. Coli	10	100	<10	Pass

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< 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	

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	

Mycotox	ins			Not	Tested
Analyte	LOQ	Limit	Mass	Status	Qualifier

Instrument: ; Method:

TLABS

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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status Qu	ıalifier	Analyte	LOQ	Limit	Mass	Status C	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		<u>.</u>
Abamectin	0.24	0.50	ND	Pass	M2	Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass		lmazalil	0.10	0.20	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	M2
Aldicarb	0.19	0.40	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass		Methyl	0.17	0.40	ND	г азз	
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	L1	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	M2	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 I1
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	1 033	
Diazinon	0.10	0.20	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	M2
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	M2
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	M2
						Trifloxystrobin	0.10	0.20	ND	Pass	

Instrument: LC-QQQ; Method: TPL_Pesticides_01



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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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