Certificate of Analysis

TRANSPARENT LABS

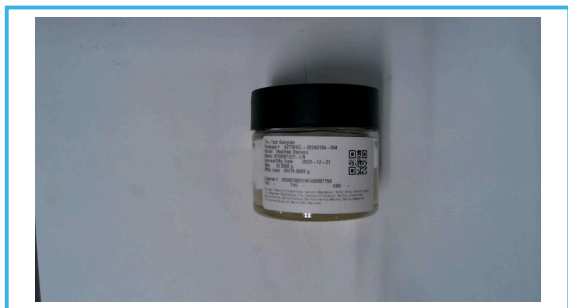## True Harvest

Phoenix, AZ 85043
mike@trueharvestco.com

Lic. #00000100DCWU00857159
Harvest Dates: 2023-12-21

Modified Banana

Concentrates & Extracts, Live Rosin, Extraction Method: Ice/Water
Dispensary License #: ; Manufacturing License #: ; Cultivation License #:

Sample: 2401TLL0006.0066
Strain: Modified Banana
Parent Batch #: ; Batch#: R1MOB1221-LR; Batch Size: 20 g
Sample Received: 01/10/2024; Report Created: 01/15/2024; Expires: 01/15/2025
Manufacturing Date:
Sampling: ; Environment:

Safety

Pass	Pass	Pass
Pesticides	Microbials	Mycotoxins
Pass	**Pass**	**Not Tested**
Solvents	Metals	Foreign Matter

Cannabinoids
TPL_Potency_01

82.82%	<LOQ	98.22%
Total THC	Total CBD	Total Cannabinoids Q3

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	93.79	937.9	M2
Δ9-THC	0.10	0.56	5.6	M2
Δ8-THC	0.10	ND	ND	M2
THCV	0.10	ND	ND	M2
CBDa	0.10	<LOQ	<LOQ	M2
CBD	0.10	ND	ND	M2
CBDV	0.10	ND	ND	M2
CBN	0.10	ND	ND	M2
CBGa	0.10	3.57	35.7	M2
CBG	0.10	0.30	3.0	M2
CBC	0.10	ND	ND	M2
Total		**98.22**	**982.2**	

Total THC = THCa * 0.877 + Δ9-THC
Total CBD = CBDa * 0.877 + CBD
Instrument: HPLC-DAD: ; Method: TPL_Potency_01

Terpenes
TPL_Terpenes_01

Hops	Lemon	Cinnamon

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
β-Myrcene		1.615	16.15	Q3
δ-Limonene		1.596	15.96	Q3
α-Humulene		1.163	11.63	Q3
β-Caryophyllene		0.968	9.68	Q3
Linalool		0.420	4.20	Q3
trans-Nerolidol		0.420	4.20	Q3
β-Pinene		0.399	3.99	Q3
α-Pinene		0.180	1.80	Q3
Ocimene		0.166	1.66	Q3
Terpinolene		0.156	1.56	Q3
γ-Terpinene		0.108	1.08	Q3
Eucalyptol		0.078	0.78	Q3
α-Bisabolol		0.077	0.77	Q3
Camphene		0.052	0.52	Q3
Caryophyllene Oxide		0.013	0.13	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		**7.411**	**74.11**	

Instrument: GCMS ; Method: TPL_Terp_01
Notes:

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Residual Solvents   Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Acetone	189.0	1000.0	ND	Pass	
Acetonitrile	77.0	410.0	ND	Pass	
Benzene	0.4	2.0	ND	Pass	
Butanes	472.0	5000.0	ND	Pass	
Chloroform	11.0	60.0	ND	Pass	
Dichloromethane	113.0	600.0	ND	Pass	
Ethanol	943.0	5000.0	ND	Pass	
Ethyl-Acetate	943.0	5000.0	ND	Pass	
Ethyl-Ether	943.0	5000.0	ND	Pass	
Heptane	943.0	5000.0	ND	Pass	
Hexanes	137.0	290.0	ND	Pass	
Isopropyl-Acetate	943.0	5000.0	ND	Pass	
Methanol	566.0	3000.0	ND	Pass	
Pentanes	943.0	5000.0	ND	Pass	
2-Propanol	943.0	5000.0	ND	Pass	
Toluene	168.0	890.0	ND	Pass	
Xylenes	819.0	2170.0	ND	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	Pass	V1
Lead	500.0	1000.0	<LOQ	Pass	
Mercury	100.0	200.0	<LOQ	Pass	

Mycotoxins   Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
B1	10	0	ND	Pass	
B2	10	0	ND	Pass	I1 M2
G1	10	0	ND	Pass	
G2	5	0	ND	Pass	
Ochratoxin A	10	20	ND	Pass	I1 L1 M1 V1
Total Aflatoxins	1	20	ND	Pass	M2

Instrument: HS-GCMS ; Method: TPL_ResSolv_01

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Sampling: ; Environment:

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	

Instrument: Plating/qPCR

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Pesticides TPL_Pesticides_01 — Pass

Analyte	LOQ PPM	Limit PPM	Mass PPM	Status	Qualifier	Analyte	LOQ PPM	Limit PPM	Mass PPM	Status	Qualifier
Abamectin	0.24	0.50	ND	Pass		Hexythiazox	0.49	1.00	ND	Pass	M2
Acephate	0.20	0.40	ND	Pass		Imazalil	0.10	0.20	ND	Pass	M2
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.20	0.40	ND	Pass	
Aldicarb	0.20	0.40	ND	Pass		Kresoxim Methyl	0.20	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenazate	0.10	0.20	ND	Pass		Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	M2	Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.20	0.40	ND	Pass		Methomyl	0.20	0.40	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	
Carbofuran	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass		Oxamyl	0.49	1.00	ND	Pass	
Chlorfenapyr	0.49	1.00	ND	Pass	I1 M2	Paclobutrazol	0.20	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2
Clofentezine	0.10	0.20	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.49	1.00	ND	Pass		Piperonyl Butoxide	0.98	2.00	ND	Pass	
Cypermethrin	0.49	1.00	ND	Pass	M2	Prallethrin	0.10	0.20	ND	Pass	
Daminozide	0.49	1.00	ND	Pass		Propiconazole	0.20	0.40	ND	Pass	
Diazinon	0.10	0.20	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Pyrethrins	0.41	1.00	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	M2
Etofenprox	0.20	0.40	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass		Spiroxamine	0.20	0.40	ND	Pass	M2
Fenpyroximate	0.20	0.40	ND	Pass		Tebuconazole	0.20	0.40	ND	Pass	
Fipronil	0.20	0.40	ND	Pass	I1	Thiacloprid	0.10	0.20	ND	Pass	
Flonicamid	0.49	1.00	ND	Pass		Thiamethoxam	0.10	0.20	ND	Pass	
Fludioxonil	0.20	0.40	ND	Pass		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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