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Total Health & Wellness dba True Harvest

Sample: 2406TLL0212.1031 Strain: Modified Banana

Parent Batch #:; Batch#: MOB240520; Batch Size: 20 g

Sample Received: 06/20/2024; Report Created: 06/26/2024; Expires: 06/26/2025

Manufacturing Date: 05/20/2024

Sampling: ; Environment:

Lic. #00000100DCWU00857159

Harvest Dates:

Phoenix, AZ 85043

Modified Banana

jpastor@trueharvestco.com

Plant, Flower - Cured, Extraction Method: Indoor

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





Safety

Pass

Pass

Pass

Pesticides

Microbials

Metals

Cannabinoids

TPL_Potency_01

30.09% Total THC <LOQ

35.65% **Total Cannabinoids** Total CBD

Analyte	LOQ	Mass	Mass	Qualifier
•	%	%	mg/g	
THCa	0.10	33.71	337.1	
Δ9-THC	0.10	0.53	5.3	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	<loq< th=""><th><loq< th=""><th></th></loq<></th></loq<>	<loq< th=""><th></th></loq<>	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	1.25	12.5	
CBG	0.10	0.17	1.7	
CBC	0.10	ND	ND	
Total		35.65	356.5	

Terpenes





LOQ	Mass	Mass	Qualifier
%	%	mg/g	
	0.1500	1.500	Q3
	0.1360	1.360	Q3
	0.1060	1.060	Q3
	0.1000	1.000	Q3
	0.0890	0.890	Q3
	0.0830	0.830	Q3
	0.0790	0.790	Q3
	0.0750	0.750	Q3
	0.0670	0.670	Q3
	0.0480	0.480	Q3
	0.0440	0.440	Q3
	0.0430	0.430	Q3
	0.0260	0.260	Q3
			Q3
			Q3
	<	<	Q3
			Q3
	<		Q3
	<	<	Q3
	1.0530	10.530	٩o
		% 0.1500 0.1360 0.1360 0.1000 0.0890 0.0830 0.0790 0.0750 0.0670 0.0440 0.0440 0.04430 0.0260 0.0040 0.0030 <	0.1500 1.500 0.1360 1.360 0.1060 1.060 0.1000 1.000 0.0890 0.890 0.0830 0.830 0.0790 0.750 0.0750 0.750 0.0670 0.670 0.0480 0.480 0.0440 0.440 0.0430 0.430 0.0260 0.260 0.0040 0.040 0.0030 0.030 <

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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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status Q	(ualifier	Analyte	LOQ	Limit	Mass	Status C	(ualifier
	PPM	PPM	PPM		<u>.</u>		PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1L1	Hexythiazox	0.48	1.00	ND	Pass	
Abamecum	0.24	0.50	ND	газэ	V1	Imazalil	0.10	0.20	ND	Pass	
Acephate	0.19	0.40	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	
Aldicarb	0.19	0.40	ND	Pass		Methyl					
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenazate	0.10	0.20	ND	Pass	V1	Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	M2	Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass		Methomyl	0.19	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	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass		Paclobutrazol	0.19	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass	M2	Permethrin	0.10	0.20	ND	Pass	M2
Clofentezine	0.10	0.20	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass	M1	Piperonyl	0.96	2.00	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1	Butoxide	0.70	2.00	ND		
Daminozide	0.48	1.00	ND	Pass		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	M2	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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Heavy Me	etals				Pass
Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	<loq< th=""><th>Pass</th><th></th></loq<>	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>V1 M1</th></loq<>	Pass	V1 M1
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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- 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.

TLABS

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