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Total Health & Wellness dba True Harvest

Sample: 2405TLL0153.0752

Strain: Modified Banana x Rotten Banana

Parent Batch #:; Batch#: MBRB240502; Batch Size: 17 g

Sample Received: 05/03/2024; Report Created: 05/10/2024; Expires: 05/10/2025 Manufacturing Date:

Sampling: ; Environment:

jpastor@trueharvestco.com

Phoenix, AZ 85043

Lic. #00000100DCWU00857159 Harvest Dates:

Modified Banana x Rotten Banana

Concentrates & Extracts, Infused/Enhanced Preroll, Extraction Method: Ice/Water Dispensary License #:; Manufacturing License #:; Cultivation License #:





Safety

Pass **Pass Pass Pesticides** Microbials **Mycotoxins Pass Pass Not Tested**

Solvents

Metals

Foreign Matter

Cannabinoids

TPL_Potency_01

36.88% <LOQ 43.40% **Total Cannabinoids** Total THC Total CBD

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	38.88	388.8	
Δ9-THC	0.10	2.79	27.9	
Δ8-ΤΗС	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	1.51	15.1	
CBG	0.10	0.22	2.2	
CBC	0.10	ND	ND	
Total		43.40	434.0	

Terpenes







Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
β-Caryophyllene		0.1600	1.600	Q3
Ocimene		0.1200	1.200	Q3
β-Myrcene		0.1100	1.100	Q3
δ-Limonene		0.1100	1.100	Q3
α-Humulene		0.0900	0.900	Q3
β-Pinene		0.0800	0.800	Q3
Terpinolene		0.0800	0.800	Q3
Eucalyptol		0.0500	0.500	Q3
trans-Nerolidol		0.0400	0.400	Q3
Linalool		0.0300	0.300	Q3
α-Bisabolol		0.0100	0.100	Q3
α-Pinene		0.0100	0.100	Q3
Camphene		0.0100	0.100	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
Caryophyllene Oxid	de	<	<	Q3
cis-Nerolidol		<	<	Q3
y-Terpinene		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		0.9000	9.000	
	•			

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 (Qualifier	Analyte	LOQ	Limit	Mass	Status (Qualifier
	PPM	PPM	PPM		<u>.</u>		PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	V1	Hexythiazox	0.48	1.00	ND	Pass	L1, V1
Acephate	0.19	0.40	ND	Pass		Imazalil	0.10	0.20	ND	Pass	M2
Acetamiprid	0.10	0.20	ND	Pass	V1	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	M1	Malathion	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	L1, V1	Metalaxyl	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	M2
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	L1, V1
Chlorantraniliprole	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	M2, V1	Oxamyl	0.48	1.00	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass	M1	Paclobutrazol	0.19	0.40	ND	Pass	
Clofentezine	0.10	0.20	ND	Pass	V1	Permethrin	0.10	0.20	ND	Pass	M2, V1
Cyfluthrin	0.48	1.00	ND	Pass	V1	Phosmet	0.10	0.20	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1, V1	Piperonyl	0.96	2.00	ND	Pass	
Daminozide	0.48	1.00	ND	Pass	M1	Butoxide	0.70	2.00	ND	Fa55	
Diazinon	0.10	0.20	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	M1, L1
Dichlorvos	0.05	0.10	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	M2
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	L1
Etofenprox	0.19	0.40	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	V1
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	
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	M2	Thiacloprid	0.10	0.20	ND	Pass	
						Thiamethoxam	0.10	0.20	ND	Pass	
						Trifloxystrobin	0.10	0.20	ND	Pass	

 $Instrument: LC\text{-}QQQ \ ; Method: TPL_Pesticides_01$



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Heavy Meta	als				Pass
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>V1</th></loq<>	Pass	V1
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>B2 L1</th></loq<>	Pass	B2 L1

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

Residual Sc	lvents				Pass
Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Acetone	476.6	1000.0	ND	Pass	
Acetonitrile	195.4	410.0	ND	Pass	
Benzene	1.0	2.0	ND	Pass	
Butanes	595.8	5000.0	ND	Pass	
Chloroform	28.6	60.0	ND	Pass	
Dichloromethane	286.0	600.0	ND	Pass	
Ethanol	2383.2	5000.0	ND	Pass	
Ethyl-Acetate	2383.2	5000.0	ND	Pass	
Ethyl-Ether	2383.2	5000.0	ND	Pass	
Heptane	2383.2	5000.0	ND	Pass	
Hexanes	138.2	290.0	ND	Pass	
Isopropyl-Acetate	2383.2	5000.0	ND	Pass	
Methanol	1429.9	3000.0	ND	Pass	
Pentanes	138.2	5000.0	ND	Pass	
2-Propanol	2383.2	5000.0	ND	Pass	
Toluene	424.2	890.0	ND	Pass	
Xvlenes	95.3	2170.0	ND	Pass	

Performed by GCMS-HS SOP-004. Methods used per AZDHS R9-17-404.03 and the solvent limits set by AZDHS R9-17 Table 3.1. AZDHS approved method for residual solvents by GCMS-HS for all listed analytes. Subcontracted through DVT Registration Certificate Identification Number: 0000031LRCHX78341676

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	

 $Instrument: qPCR/Plating; AOAC\ Methods\ 082102, 022202\ and\ 2018.13$

Mycotoxins					Pass
Analyte	LOQ	Limit	Mass	StatusC	Qualifier
	PPB	PPB	PPB		
B1	8.1	20.0	ND	Pass	M2
B2	8.1	20.0	ND	Pass	M2
G1	8.1	20.0	ND	Pass	M2 L1
G2	8.1	20.0	ND	Pass	L1
Ochratoxin A	8.1	20.0	ND	Pass	V1L1
Total Aflatoxins	8.1	20.0	ND	Pass	L1 M2



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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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