

Amended

Powered by Confident LIMS



Strain: Melonade

Parent Batch #: ; Batch#: R4MEL0312; Batch Size: 20 g

Sample Received: 04/01/2024; Report Created: 05/10/2024; Expires: 05/10/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159

jpastor@trueharvestco.com

Harvest Dates: 03/12/2024 **Melonade**

Phoenix, AZ 85043

Plant, Flower - Cured

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

Total Health & Wellness dba True Harvest





Safety

Pass

Pass

Pass

Pesticides

Microbials

Metals

Cannabinoids

TPL_Potency_01

Analyte
THCa
Δ9-THC
Δ8-THC

THCV

CBDa

CBD CBDV

CBN CBGa

CBG

CBC

Total

25.08%Total THC

<LOQ
Total CBD

LOO

0.10

0.10

0.10

0.10

0.10

0.10

0.10

0.12

ND

29.93

29.93% Total Cannabinoids O3

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Mass	Mass	Qualifier
%	mg/g	
28.33	283.3	
0.23	2.3	
ND	ND	
ND	ND	
LOQ	<loq< td=""><td></td></loq<>	
ND	ND	
ND	ND	
ND	ND	
1.25	12.5	

1.2

ND

299.3

Terpenes



Lemon





Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
δ-Limonene		0.4000	4.000	Q3
β-Caryophyllene		0.2500	2.500	Q3
α-Humulene		0.2300	2.300	Q3
β-Myrcene		0.1400	1.400	Q3
β-Pinene		0.1200	1.200	Q3
Guaiol		0.1000	1.000	Q3
Ocimene		0.1000	1.000	Q3
trans-Nerolidol		0.1000	1.000	Q3
Linalool		0.0900	0.900	Q3
Terpinolene		0.0800	0.800	Q3
y-Terpinene		0.0700	0.700	Q3
α-Bisabolol		0.0600	0.600	Q3
Eucalyptol		0.0500	0.500	Q3
α-Pinene		0.0400	0.400	Q3
Camphene		0.0100	0.100	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
Caryophyllene Oxide		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		1.8400	18.400	-

Instrument: GCMS; Method: TPL_Terp_01

Notes

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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Total Health & Wellness dba True Harvest



Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status C	Qualifier	Analyte	LOQ	Limit	Mass	Status C	(ualifier
	PPM	PPM	PPM				PPM	PPM	PPM		<u> </u>
Abamectin	0.24	0.50	ND	Pass	M1 R2	Hexythiazox	0.48	1.00	ND	Pass	M2
Abamecum	0.24	0.50	ND	F a 5 5	L1 I1	lmazalil	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	0.17	0.40			
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	L1 R2
Bifenazate	0.10	0.20	ND	Pass		Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass		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	V1
Chlorantraniliprole	0.10	0.20	ND	Pass		Oxamyl	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	M2	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		Piperonyl	0.96	2.00	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1	Butoxide	0.70	2.00	ND	1 033	
Daminozide	0.48	1.00	ND	Pass	M1 R2	Prallethrin	0.10	0.20	ND	Pass	M1
Daminozide					L1	Propiconazole	0.19	0.40	ND	Pass	M2
Diazinon	0.10	0.20	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Pyrethrins	0.48	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	
Etofenprox	0.19	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	0.19	0.40	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass		Thiacloprid	0.10	0.20	ND	Pass	
Flonicamid	0.48	1.00	ND	Pass		Thiamethoxam	0.10	0.20	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass		Trifloxystrobin	0.10	0.20	ND	Pass	

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



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Powered by Confident LIMS 3 of 4

Total Health & Wellness dba True Harvest Sample: 2404TLL0111.0575

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Heavy Metals					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></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	<u> </u>
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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Amended: Client requested that the following harvest date be added: 03/12/2024

- 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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