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Total Health & Wellness dba True Harvest

Sample: 2405TLL0173.0841

Strain: Wedding Pie

Parent Batch #:; Batch#: WDP240429; Batch Size: 20 g

Sample Received: 05/20/2024; Report Created: 05/21/2024; Expires: 05/21/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159

Harvest Dates:

Wedding Pie

Phoenix, AZ 85043 jpastor@trueharvestco.com

Plant, Flower - Cured

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



Safety

Pass

Pass

Pass

Pesticides

Microbials

Metals

Cannabinoids

TPL_Potency_01

28.71% Total THC <LOQ

33.79% **Total Cannabinoids** Total CBD

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	32.21	322.1	
Δ9-ΤΗС	0.10	0.46	4.6	
Δ8-ΤΗС	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	0.98	9.8	
CBG	0.10	0.14	1.4	
CBC	0.10	ND	ND	
Total		33.79	337.9	

Terpenes TPL_Terpenes_01







Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
β-Caryophyllene		0.2500	2.500	Q3
Linalool		0.2400	2.400	Q3
α-Humulene		0.2200	2.200	Q3
δ-Limonene		0.1700	1.700	Q3
Ocimene		0.1700	1.700	Q3
Guaiol		0.1400	1.400	Q3
β-Myrcene		0.1300	1.300	Q3
trans-Nerolidol		0.1200	1.200	Q3
β-Pinene		0.0900	0.900	Q3
Terpinolene		0.0800	0.800	Q3
y-Terpinene		0.0700	0.700	Q3
α-Bisabolol		0.0500	0.500	Q3
Eucalyptol		0.0400	0.400	Q3
α-Pinene		0.0200	0.200	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

Instrument: GCMS; Method: TPL_Terp_01

Total

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 Q	ualifier
	PPM	PPM	PPM		<u> </u>		PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	V1	Hexythiazox	0.48	1.00	ND	Pass	M1
Acephate	0.19	0.40	ND	Pass		lmazalil	0.10	0.20	ND	Pass	M1
Acetamiprid	0.10	0.20	ND	Pass	M1	Imidacloprid	0.19	0.40	ND	Pass	M1
Aldicarb	0.19	0.40	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	М1
Azoxystrobin	0.10	0.20	ND	Pass	M1	Methyl	0.17	0.40	ND	F 455	1417
Bifenazate	0.10	0.20	ND	Pass	M1L1	Malathion	0.10	0.20	ND	Pass	M1
Direitazate	0.10			1 433	V1	Metalaxyl	0.10	0.20	ND	Pass	M1
Bifenthrin	0.10	0.20	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass	M1	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	M1	Oxamyl	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	M1 V1	Paclobutrazol	0.19	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	0.48	1.00	ND	Pass		Piperonyl	0.96	2.00	ND	Pass	М1
Cypermethrin	0.48	1.00	ND	Pass	M1	Butoxide					
Daminozide	0.48	1.00	ND	Pass	M1L1	Prallethrin	0.10	0.20	ND	Pass	
					V1	Propiconazole	0.19	0.40	ND	Pass	M1
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	M1
Ethoprophos	0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	M1
Etofenprox	0.19	0.40	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	M1
Etoxazole	0.10	0.20	ND	Pass	M1	Spirotetramat	0.10	0.20	ND	Pass	M1
Fenoxycarb	0.10	0.20	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	M1
Fenpyroximate	0.19	0.40	ND	Pass	M1	Tebuconazole	0.19	0.40	ND	Pass	M1
Fipronil	0.19	0.40	ND	Pass		Thiacloprid	0.10	0.20	ND	Pass	M1
Flonicamid	0.48	1.00	ND	Pass	M1	Thiamethoxam	0.10	0.20	ND	Pass	M1
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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Heavy Me	tals				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	ND	Pass	
Morcury	1000	200.0	<100	Dacc	

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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- 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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