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### Total Health & Wellness dba True Harvest

Sample: 2404TLL0150.0731

Strain: Frosted Donuts

Parent Batch #:; Batch#: R2FD0410; Batch Size: 20 g

Sample Received: 04/29/2024; Report Created: 05/03/2024; Expires: 05/03/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159

Harvest Dates:

#### **Frosted Donuts**

jpastor@trueharvestco.com

Plant, Flower - Cured

Phoenix, AZ 85043

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





#### Safety

**Pass** 

**Pass** 

**Pass** 

**Pesticides** 

Microbials

Metals

Cannabinoids

TPL\_Potency\_01

25.21% **Total THC** 

ND Total CBD

29.80% **Total Cannabinoids** 

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	28.40	284.0	
Δ9-THC	0.10	0.30	3.0	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	ND	ND	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	0.93	9.3	
CBG	0.10	0.17	1.7	
CBC	0.10	ND	ND	
Total		29 80	298.0	

Terpenes TPL\_Terpenes\_01



Lemon





Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	<u> </u>
δ-Limonene		0.3570	3.570	Q3
Ocimene		0.3220	3.220	Q3
Linalool		0.2450	2.450	Q3
β-Caryophyllene		0.1920	1.920	Q3
α-Humulene		0.1420	1.420	Q3
β-Myrcene		0.1320	1.320	Q3
β-Pinene		0.1260	1.260	Q3
trans-Nerolidol		0.1030	1.030	Q3
Terpinolene		0.0770	0.770	Q3
y-Terpinene		0.0670	0.670	Q3
α-Terpinene		0.0610	0.610	Q3
α-Pinene		0.0520	0.520	Q3
Eucalyptol		0.0440	0.440	Q3
Camphene		0.0130	0.130	Q3 Q3
3-Carene		<	<	Q3
α-Bisabolol		<	<	Q3
Caryophyllene Oxide		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		1.9330	19.330	

Instrument: GCMS; Method: TPL\_Terp\_01

Total THC = THCa \* 0.877 +  $\Delta$ 9-THC Total CBD = CBDa \* 0.877 + CBD Instrument: HPLC-DAD: ; Method: TPL\_Potency\_01

LABS

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Brian DiMarco **Laboratory Director** 

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### Pesticides TPL\_Pesticides\_01

**Pass** 

Analyte	LOQ	Limit	Mass	Status C	Qualifier	Analyte	LOQ	Limit	Mass	Status Q	ualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1L1	Hexythiazox	0.48	1.00	ND	Pass	M2
Abameetin					R2 V1	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					
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		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	M2	Paclobutrazol	0.19	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	
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	M1L1	Butoxide					
Daminozide	0.48	1.00	ND	Pass	M1 V1	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	M4
Ethoprophos	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass		Spinosad	0.10	0.20	ND	Pass	M2
Etoxazole	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass	M2	Spirotetramat	0.10	0.20	ND	Pass	M1
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	M2
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	M2

 $Instrument: LC\text{-}QQQ \ ; Method: TPL\_Pesticides\_01$ 



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Heavy Me	etais				Pass
Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	
Cadmium	200.0	400.0	<loq< td=""><td>Pass</td><td></td></loq<>	Pass	
Lead	500.0	1000.0	<loq< td=""><td>Pass</td><td></td></loq<>	Pass	
M =	1000	200.0	400	D	

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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