

Powered by Confident LIMS 1 of 4

Total Health & Wellness dba True Harvest

Sample: 2405TLL0173.0840

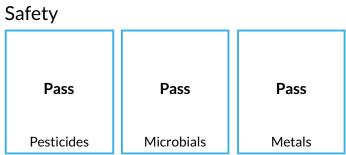
Phoenix, AZ 85043 jpastor@trueharvestco.com

Strain: Frosted Donuts Parent Batch #: ; Batch #: FRD240429; 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: **Frosted Donuts**

Plant, Flower - Cured Dispensary License #: ; Manufacturing License #: ; Cultivation License #:





Cannabinoids

	_		
<loq< th=""><th></th><th>31.56</th><th>%</th></loq<>		31.56	%
Total CBD		Total Canna Q3	binoids
LOQ	Mass	Mass	Qualifier
%	%	mg/g	
0.10	29.97	299.7	
0.10	0.40	4.0	
0.10	ND	ND	
0.10	ND	ND	
0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
0.10	ND	ND	
0.10	ND	ND	
0.10	ND	ND	
0.10	1.00	10.0	
0.10	0.19	1.9	
0.10	ND	ND	
	31.56	315.6	
	Kod 100 % 0.10	Total CBD LOQ Mass % % 0.10 29.97 0.10 0.40 0.10 ND 0.10 ND	Total CBD Total Canaga LOQ Mass Mass % % mg/g 0.10 29.97 299.7 0.10 0.40 4.0 0.10 0.40 4.0 0.10 ND ND 0.10 0.19 1.9 0.10 ND ND

Total THC = THCa * $0.877 + \Delta 9$ -THC Total CBD = CBDa * 0.877 + CBDInstrument: HPLC-DAD: ; Method: TPL_Potency_01



1721 E McDowell Road Phoenix, AZ (602) 368-4233 transparentlabsaz.com Lic#0000029LRCXG19240160 Instrument: GCMS; Method: TPL_Terp_01 Notes:

Brian DiMarco

Laboratory Director

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Ternenes

TPL_Terpenes_01				
<u>k</u>	8		1	
Lavender	Lemon		Ear	thy
Analyte	LOQ	Mass	Mass	Qualifier
$ \begin{array}{ c c c c } Linalool & & & \\ \hline \delta-Limonene & & \\ Ocimene & & \\ \hline Ocimene & & \\ \hline & \\ \partial-Limone & & \\ \hline & \\ \beta-Pinene & & \\ \hline & \\ Terpinolene & & \\ Trapinene & & \\ Linal & \\ \hline & \\ v-Terpinene & \\ Caryophyllone & & \\ \hline & \\ \alpha-Pinene & & \\ \hline &$	%	% 0.2100 0.2000 0.2000 0.1900 0.1900 0.1000 0.0900 0.0900 0.0900 0.0400 0.0300 0.0400 0.0300 0.0100 < < < <	mg/g 2.100 2.000 2.000 1.900 1.300 1.000 0.900 0.900 0.900 0.900 0.700 0.400 0.300 0.100 < < < < < < < < < < 14.600	23 23 23 23 23 23 23 23 23 23 23 23 23 2
Instrument: CCMS: Method:	TPI Tern 01			



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Pesticides TPL_Pesticides_01

Analyte	LOQ	Limit	Mass	Status C	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		Imazalil	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	M1
Azoxystrobin	0.10	0.20	ND	Pass	M1	Methyl	0.19	0.40	ND	PdSS	IVII
Bifenazate	0.10	0.20	ND	Pass	M1L1	Malathion	0.10	0.20	ND	Pass	M1
Diferiazate	0.10	0.20	ND	F d 5 5	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	M1
Cypermethrin	0.48	1.00	ND	Pass	M1	Butoxide					1411
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-QQQ ; Method: TPL_Pesticides_01



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Heavy Metals

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	
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,

11 = 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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