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Total Health & Wellness dba True Harvest

Sample: 2403TLL0080.0416

Phoenix, AZ 85043 jpastor@trueharvestco.com

Strain: Red Velvet Parent Batch #:; Batch#: R3JSR0122; Batch Size: 20 g Sample Received: 03/06/2024; Report Created: 03/12/2024; Expires: 03/12/2025 Manufacturing Date: Sampling: ; Environment:

Lic. #00000100DCWU00857159 Harvest Dates: Red Velvet

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



Safety Pass Pass Pass Pesticides Microbials Metals

Cannabinoids

1	PL_Potency_01						
	26.24%		<loq< th=""><th></th><th></th><th>30.66</th><th>%</th></loq<>			30.66	%
	Total THC	Т	otal CBD			Total Canna Q3	binoids
Analyte			LOQ	Mass		Mass	Qualifier
			%		%	mg/g	
Т	HCa		0.10	29.3	88	293.8	
Δ	9-THC		0.10	0.4	18	4.8	
Δ	8-THC		0.10	N	D	ND	
Т	HCV		0.10	N	D	ND	
С	BDa		0.10	N	D	ND	
С	BD		0.10	N	D	ND	
С	BDV		0.10	N	D	ND	
С	BN		0.10	Ν	D	ND	
С	BGa		0.10	0.8	31	8.1	
С	BG		0.10	Ν	D	ND	
CBC			0.10	ND		ND	
Total				30.6	66	306.6	

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



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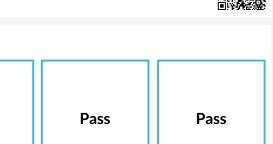
Instrument: GCMS; Method: TPL_Terp_01 Notes: hiti

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Terpenes

IPL_Ierpenes_01				
<u>k</u>	Ý		ų	È
Lavender	Cinnamon		Ho	ops
Analyte	LOQ	Mass	Mass	Qualifier
Linalool β-Caryophyllene α-Humulene δ-Limonene trans-Nerolidol β-Pinene β-Myrcene Ocimene Terpinolene	%	% 0.5600 0.2800 0.2700 0.2700 0.2300 0.1300 0.1200 0.0900 0.0800	mg/g 5.600 2.800 2.700 2.700 2.300 1.300 1.200 0.900 0.800	23 23 23 23 23 23 23 23 23 23 23 23
y-Terpinene Eucalyptol α-Bisabolol α-Pinene Camphene Caryophyllene Oxide 3-Carene α-Terpinene cis-Nerolidol Geraniol Guaiol Isopulegol		0.0700 0.0500 0.0400 0.0200 0.0100 < < < < < < < < < < < <	0.700 0.500 0.400 0.200 0.100 < < < < < < < < < <	23 23 23 23 23 23 23 23 23 23 23 23 23 2
p-Ċymene Total		< 2.2600	< 22.600	Q3 Q3
Instrument: GCMS: Method	: TPL Terp 01			



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Pass

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Pesticides TPL_Pesticides_01

Analyte	LOQ	Limit	Mass	Status C	Qualifier	Analyte	LOQ	Limit	Mass	Status C	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1L1	Hexythiazox	0.48	1.00	ND	Pass	V1
Abamectin	0.24	0.50	ND	Fd55	V1	Imazalil	0.10	0.20	ND	Pass	
Acephate	0.19	0.40	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	V1
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	V1	Methiocarb	0.10	0.20	ND	Pass	L1
Boscalid	0.19	0.40	ND	Pass		Methomyl	0.19	0.40	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass	L1	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	M1L1	Oxamyl	0.48	1.00	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	V1	Paclobutrazol	0.19	0.40	ND	Pass	V1
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2 V1
Clofentezine	0.10	0.20	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass	M1V1	Piperonyl	0.96	2.00	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1V1	Butoxide				1 455	
Daminozide	0.48	1.00	ND	Pass	M1L1	Prallethrin	0.10	0.20	ND	Pass	M1V1
Diazinon	0.10	0.20	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass	M1V1	Propoxur	0.10	0.20	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Pyrethrins	0.48	1.00	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	V1
Etofenprox	0.19	0.40	ND	Pass		Spinosad	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenoxycarb	0.10	0.20	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	V1
Fenpyroximate	0.19	0.40	ND	Pass	V1	Spiroxamine	0.19	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass	V1	Tebuconazole	0.19	0.40	ND	Pass	V1
Flonicamid	0.48	1.00	ND	Pass		Thiacloprid	0.10	0.20	ND	Pass	V1
Fludioxonil	0.19	0.40	ND	Pass		Thiamethoxam	0.10	0.20	ND	Pass	V1
						Trifloxystrobin	0.10	0.20	ND	Pass	

Instrument: LC-QQQ ; Method: TPL_Pesticides_01



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Pass

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Heavy Metals

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	<loq< th=""><th>Pass</th><th></th></loq<>	Pass	
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></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	

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 (0)

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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