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Total Health & Wellness dba True Harvest

Sample: 2405TLL0160.0791 Strain: Cadillac Rainbows

Parent Batch #:; Batch#: R6CAD0417; Batch Size: 20 g

Sample Received: 05/08/2024; Report Created: 05/15/2024; Expires: 05/15/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159

Harvest Dates:

Cadillac Rainbow

Plant, Flower - Cured

Phoenix, AZ 85043 jpastor@trueharvestco.com

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





Safety

Pass

Pass

Pass

Pesticides

Microbials

Metals

Cannabinoids

TPL_Potency_01

32.95% **Total THC**

<LOQ Total CBD

38.91% **Total Cannabinoids**

Analyte	LOQ	Mass	Mass	Qualifie
	%	%	mg/g	
THCa	0.10	37.21	372.1	
Δ9-THC	0.10	0.31	3.1	
Δ8-ΤΗС	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	<loq< td=""><td><loq< td=""><td></td></loq<></td></loq<>	<loq< td=""><td></td></loq<>	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	1.26	12.6	
CBG	0.10	0.13	1.3	
CBC	0.10	ND	ND	
Total		38.91	389.1	

Terpenes TPL_Terpenes_01



Instrument: GCMS; Method: TPL_Terp_01





Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
β-Caryophyllene		0.2600	2.600	Q3
α-Humulene		0.2400	2.400	Q3
δ-Limonene		0.2300	2.300	Q3
Ocimene		0.2100	2.100	Q3
β-Myrcene		0.1500	1.500	Q3
β-Pinene		0.1100	1.100	Q3
Linalool		0.1100	1.100	Q3
Terpinolene		0.0800	0.800	Q3
trans-Nerolidol		0.0800	0.800	Q3
y-Terpinene		0.0700	0.700	Q3
Eucalyptol		0.0500	0.500	Q3
α-Pinene		0.0300	0.300	Q3
α-Bisabolol		0.0100	0.100	Q3
Camphene		0.0100	0.100	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
Caryophyllene Oxide		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		1.6400	16.400	

Total THC = THCa * 0.877 + Δ 9-THC Total CBD = CBDa * 0.877 + CBD Instrument: HPLC-DAD: ; Method: TPL_Potency_01

LABS

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Phoenix, AZ (602) 368-4233

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Brian DiMarco **Laboratory Director**

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Sample: 2405TLL0160.0791

Strain: Cadillac Rainbows

Parent Batch #:; Batch#: R6CAD0417; Batch Size: 20 g

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Pesticides TPL_Pesticides_01

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	M2	Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass		lmazalil	0.10	0.20	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
Aldicarb	0.19	0.40	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	V1
Azoxystrobin	0.10	0.20	ND	Pass		Methyl	0.17	0.40		F d 3 3	
Bifenazate	0.10	0.20	ND	Pass	M1L1	Malathion	0.10	0.20	ND	Pass	L1 R1
					V1	Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	M2
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	V1	Paclobutrazol	0.19	0.40	ND	Pass	M2
Chlorpyrifos	0.10	0.20	ND	Pass	M2	Permethrin	0.10	0.20	ND	Pass	M2
Clofentezine	0.10	0.20	ND	Pass	M2	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	M2 R1	Butoxide	0.70	2.00	ND	1 033	
Daminozide	0.48	1.00	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	M2
Diazinon	0.10	0.20	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass	M1L1	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	
Etofenprox	0.19	0.40	ND	Pass		Spinosad	0.10	0.20	ND	Pass	L1
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		Spiroxamine	0.19	0.40	ND	Pass	
Fipronil	0.19	0.40	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
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	

Instrument: LC-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	V1
Cadmium	200.0	400.0	<loq< th=""><th>Pass</th><th>V1</th></loq<>	Pass	V1
Lead	500.0	1000.0	ND	Pass	
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>V1 L1</th></loq<>	Pass	V1 L1

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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