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#### Total Health & Wellness dba True Harvest

Sample: 2403TLL0106.0540

Strain: Orion's Belt

Phoenix, AZ 85043 Parent Batch #:; Batch#: R5OB0304; Batch Size: 20 g jpastor@trueharvestco.com

Sample Received: 03/27/2024; Report Created: 04/02/2024; Expires: 04/02/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159 Harvest Dates:

Orion's Belt

Plant, Flower - Cured

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





#### Safety

**Pass** 

**Pass** 

**Pass** 

**Pesticides** 

Microbials

Metals

Cannabinoids

TPL\_Potency\_01

25.84% Total THC

ND Total CBD

30.43% **Total Cannabinoids** 

Analyte	LOQ	Mass	Mass	Qualifie
	%	%	mg/g	
THCa	0.10	29.14	291.4	
Δ9-THC	0.10	0.29	2.9	
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	0.79	7.9	
CBG	0.10	0.21	2.1	
CBC	0.10	ND	ND	
Total		30.43	304.3	

Terpenes TPL\_Terpenes\_01



Hops







Qualifier Analyte Mass Mass 0.6200 mg/g 6.200 α-Humulene 0.5400 5.400 **β-Caryophyllene** δ-Limonene 0.2300 2.300 Ocimene 0.2200 2.200 0.1500 1.500 Linalool 1.400 trans-Nerolidol 0.1400 **β-Myrcene** 0.1200 1.200 B-Pinene 0.1100 1.100 α-Bisabolol 0.0900 0.900 Terpinolene 0.0800 0.800 y-Terpinene 0.0700 0.700 α-Pinene 0.0500 0.500 0.0500 0.500 Eucalyptol 0.0100 0.100 Camphene Caryophyllene Oxide 0.0100 0.100 3-Carene α-Terpinene cis-Nerolidol Geraniol Guaiol Isopulegol p-Cymene Total 2.4900

Instrument: GCMS; Method: TPL\_Terp\_01

Total THC = THCa \* 0.877 + Δ9-THC Total CBD = CBDa \* 0.877 + CBD

Instrument: HPLC-DAD: ; Method: TPL\_Potency\_01

LABS

1721 E McDowell Road Phoenix, AZ (602) 368-4233 transparentlabsaz.com Lic#0000029LRCXG19240160

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	V1	Hexythiazox	0.48	1.00	ND	Pass	M2
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	
Azoxystrobin	0.10	0.20	ND	Pass		Methyl	0.17	0.40	ND	Fa55	
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass		Metalaxyl	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Methomyl	0.19	0.40	ND	Pass	
Carbofuran	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorfenapyr	0.48	1.00	ND	Pass	M1	Oxamyl	0.48	1.00	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass	M2	Paclobutrazol	0.19	0.40	ND	Pass	
Clofentezine	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2
Cyfluthrin	0.48	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1	Piperonyl	0.96	2.00	ND	Pass	
Daminozide	0.48	1.00	ND	Pass	M1L1	Butoxide	0.70	2.00	ND	F455	
Diazinon	0.10	0.20	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Propiconazole	0.19	0.40	ND	Pass	
Dimethoate	0.10	0.20	ND	Pass		Propoxur	0.10	0.20	ND	Pass	
Ethoprophos	0.10	0.20	ND	Pass		Pyrethrins	0.48	1.00	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass		Pyridaben	0.10	0.20	ND	Pass	
Etoxazole	0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	M2
Fenoxycarb	0.10	0.20	ND	Pass		Spiromesifen	0.10	0.20	ND	Pass	
Fenpyroximate	0.19	0.40	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	M1
Fipronil	0.19	0.40	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	
Flonicamid	0.48	1.00	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
Fludioxonil	0.19	0.40	ND	Pass		Thiacloprid	0.10	0.20	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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man

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Heavy Me	etals				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	<loq< th=""><th>Pass</th><th>V1</th></loq<>	Pass	V1
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>V1</th></loq<>	Pass	V1

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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