

True Harvest

Sample: 2401TLL0006.0062

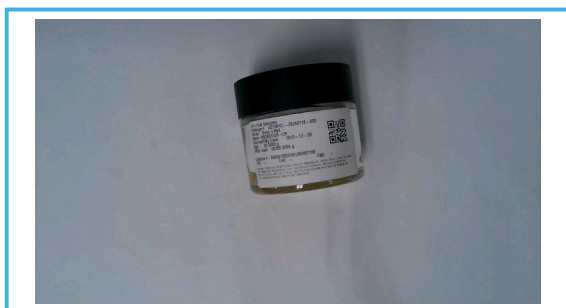
Phoenix, AZ 85043
mike@trueharvestco.com

Strain: Orion's Belt
Parent Batch #: ; Batch#: R5OB01228-LR; Batch Size: 20 undefined
Sample Received: 01/10/2024; Report Created: 01/15/2024; Expires: 01/15/2025
Manufacturing Date:
Sampling: ; Environment:

Lic. #00000100DCWU00857159
Harvest Dates: 2023-12-28

Orion's Belt

Concentrates & Extracts, Live Rosin, Extraction Method: Ice/Water
Dispensary License #: ; Manufacturing License #: ; Cultivation License #:



Safety

Pass Pesticides	Pass Microbials	Pass Mycotoxins
Pass Solvents	Pass Metals	Not Tested Foreign Matter

Cannabinoids

TPL_Potency_01

81.20%	0.11%	95.58%
Total THC	Total CBD	Total Cannabinoids Q3

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	92.18	921.8	
Δ9-THC	0.10	0.36	3.6	
Δ8-THC	0.10	ND	ND	
THCV	0.10	ND	ND	
CBDa	0.10	0.12	1.2	
CBD	0.10	ND	ND	
CBDV	0.10	ND	ND	
CBN	0.10	ND	ND	
CBGa	0.10	2.44	24.4	
CBG	0.10	0.48	4.8	
CBC	0.10	ND	ND	
Total		95.58	955.8	

Total THC = THCa * 0.877 + Δ9-THC
Total CBD = CBDa * 0.877 + CBD
Instrument: HPLC-DAD: ; Method: TPL_Potency_01

Terpenes

TPL_Terpenes_01

Hops	Cinnamon	Lemon
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Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
α-Humulene		1.378	13.78	Q3
β-Caryophyllene		1.162	11.62	Q3
δ-Limonene		0.613	6.13	Q3
Ocimene		0.557	5.57	Q3
trans-Nerolidol		0.352	3.52	Q3
Linalool		0.312	3.12	Q3
α-Bisabolol		0.212	2.12	Q3
β-Pinene		0.195	1.95	Q3
α-Pinene		0.162	1.62	Q3
β-Myrcene		0.137	1.37	Q3
Guaiol		0.078	0.78	Q3
Terpinolene		0.066	0.66	Q3
γ-Terpinene		0.054	0.54	Q3
Eucalyptol		0.041	0.41	Q3
Camphene		0.029	0.29	Q3
Caryophyllene Oxide		0.017	0.17	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Total		5.365	53.65	

Instrument: GCMS; Method: TPL_Terp_01
Notes:

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Residual Solvents

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM		
Acetone	192.0	1000.0	ND	Pass	
Acetonitrile	79.0	410.0	ND	Pass	
Benzene	0.4	2.0	ND	Pass	
Butanes	481.0	5000.0	ND	Pass	
Chloroform	12.0	60.0	ND	Pass	
Dichloromethane	115.0	600.0	ND	Pass	
Ethanol	962.0	5000.0	ND	Pass	
Ethyl-Acetate	962.0	5000.0	ND	Pass	
Ethyl-Ether	962.0	5000.0	ND	Pass	
Heptane	962.0	5000.0	ND	Pass	
Hexanes	139.0	290.0	ND	Pass	
Isopropyl-Acetate	962.0	5000.0	ND	Pass	
Methanol	577.0	3000.0	ND	Pass	
Pentanes	962.0	5000.0	ND	Pass	
2-Propanol	962.0	5000.0	ND	Pass	
Toluene	171.0	890.0	ND	Pass	
Xylenes	835.0	2170.0	ND	Pass	

Heavy Metals

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	Pass	
Cadmium	200.0	400.0	<LOQ	Pass	V1
Lead	500.0	1000.0	<LOQ	Pass	
Mercury	100.0	200.0	<LOQ	Pass	

Mycotoxins

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
B1	10	0	ND	Pass	
B2	10	0	ND	Pass	I1 M2
G1	10	0	ND	Pass	
G2	5	0	ND	Pass	
Ochratoxin A	10	20	ND	Pass	I1 L1 M1 V1
Total Aflatoxins	1	20	ND	Pass	M2

Instrument: HS-GCMS ; Method: TPL_ResSolv_01

1721 E McDowell Road
Phoenix, AZ
(631) 953-6921
<https://www.transparentlabsaz.com>
Lic# 0000029LRCXG19240160

Brian DiMarco
Laboratory Director

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coa.support@confidentlims.com
(866) 506-5866
www.confidentlims.com





Certificate of Analysis

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Microbials

Pass

Analyte	LOQ	Limit	Result	Status	Qualifier
E. Coli	CFU/g 10	CFU/g 100	CFU/g <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: qPCR/Plating

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coa.support@confidentlims.com
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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status	Qualifier	Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.25	0.50	ND	Pass		Hexythiazox	0.50	1.00	ND	Pass	M2
Acephate	0.20	0.40	ND	Pass		Imazalil	0.10	0.20	ND	Pass	M2
Acetamiprid	0.10	0.20	ND	Pass		Imidacloprid	0.20	0.40	ND	Pass	
Aldicarb	0.20	0.40	ND	Pass		Kresoxim	0.20	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass		Methyl					
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	M2	Metalaxyl	0.10	0.20	ND	Pass	
Boscalid	0.20	0.40	ND	Pass		Methiocarb	0.10	0.20	ND	Pass	
Carbaryl	0.10	0.20	ND	Pass		Methomyl	0.20	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	0.25	0.50	ND	Pass	
Chlorfenapyr	0.50	1.00	ND	Pass	I1 M2	Oxamyl	0.50	1.00	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Paclobutrazol	0.20	0.40	ND	Pass	
Clofentezine	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2
Cyfluthrin	0.50	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cypermethrin	0.50	1.00	ND	Pass	M2	Piperonyl	0.99	2.00	ND	Pass	
Daminozide	0.50	1.00	ND	Pass		Butoxide					
Diazinon	0.10	0.20	ND	Pass		Prallethrin	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass		Propiconazole	0.20	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	0.42	1.00	ND	Pass	
Etofenprox	0.20	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	0.20	0.40	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
Fipronil	0.20	0.40	ND	Pass	I1	Spiroxamine	0.20	0.40	ND	Pass	M2
Flonicamid	0.50	1.00	ND	Pass		Tebuconazole	0.20	0.40	ND	Pass	
Fludioxonil	0.20	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	

Instrument: LC-QQQ ; Method: TPL_Pesticides_01

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B1 = Target analyte detected in calibration blank was above LOQ but the concentration of cannabinoid was below 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(I)(1)(b), but the sample's target analytes were not detected above the maximum allowable concentrations for the analytes in the sample.