

Amended

Powered by Confident LIMS

Sample: 2407TLL0250.1211

Strain: Headhunter

Parent Batch #:; Batch#: 32071024TM-HH; Batch Size: g Sample Received: 07/19/2024; Report Created: 09/16/2024

Manufacturing Date:

Sampling: ; Environment:

### High Grade Confidential

Phoenix, AZ 85042

Lic. #00000077DCPS00216601 Harvest Dates: 07/10/24

Lape' Farms: Seaside

Plant, Other, Extraction Method: Indoor

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



### Safety

Pass

Pesticides

**Pass** 

Microbials

**Pass** 

Metals

### Cannabinoids

TPL\_Potency\_01

29.16% Total THC <LOQ
Total CBD

**34.34%**Total Cannabinoids
Q3

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
THCa	0.10	32.64	326.4	
Δ9-ΤΗС	0.10	0.53	5.3	
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	1.04	10.4	
CBG	0.10	0.14	1.4	
CBC	0.10	ND	ND	
Total		34.34	343.4	

Terpenes TPL\_Terpenes\_01



Cinnamon

\*

Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
δ-Limonene		0.2120	2.120	Q3
β-Caryophyllene		0.2070	2.070	Q3
α-Humulene		0.1980	1.980	Q3
Linalool		0.1310	1.310	Q3
β-Pinene		0.0570	0.570	Q3
α-Pinene		0.0540	0.540	Q3
trans-Nerolidol		0.0480	0.480	Q3
β-Myrcene		0.0250	0.250	Q3
Ocimene		0.0100	0.100	Q3
Camphene		0.0060	0.060	Q3
3-Carene		<	<	Q3
α-Bisabolol		<	<	Q3
α-Terpinene		<	<	Q3
Caryophyllene Oxide		<	<	Q3
cis-Nerolidol		<	<	Q3
Eucalyptol		<	<	Q3
y-Terpinene		<	<	Q3
Geraniol		<	<	Q3
Guaiol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3
Terpinolene		<	<	Q3
Total		0.9480	9.480	

Instrument: GCMS; Method: TPL\_Terp\_01 Notes:

Total THC = THCa  $^*$  0.877 +  $\Delta$ 9-THC Total CBD = CBDa  $^*$  0.877 + CBD Instrument: HPLC-DAD: ; Method: TPL\_Potency\_01

TLABS

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hipin

Brian DiMarco Laboratory Director





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### Pesticides TPL\_Pesticides\_01

**Pass** 

Analyte	LOQ	Limit	Result	Status Qu	ıalifier	Analyte	LOQ	Limit	Result	Status Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM	
Abamectin	0.24	0.50	ND	Pass	R1	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
Azoxystrobin	0.10	0.20	ND	Pass		Methyl	0.19	0.40	ND	Pass
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		Oxamyl	0.48	1.00	ND	Pass
Chlorpyrifos	0.10	0.20	ND	Pass		Paclobutrazol	0.19	0.40	ND	Pass
Clofentezine	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass
Cyfluthrin	0.48	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass
Cypermethrin	0.48	1.00	ND	Pass		Piperonyl	0.96	2.00	ND	Pass
Daminozide	0.48	1.00	ND	Pass	R1	Butoxide	0.70	2.00	ND	Fd55
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
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
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

 $Instrument: LC\text{-}QQQ \ ; Method: TPL\_Pesticides\_01$ 



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Brian DiMarco

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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></th></loq<>	Pass	
Lead	500.0	1000.0	<loq< th=""><th>Pass</th><th></th></loq<>	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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**Amended** 

Amended: Sample name changed from "Headhunter Flower" to "Lape' Farms - Seaside".

- 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.



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