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SROAZ LLC

Tempe, AZ 85281

Sample: 2404TLL0133.0657

Strain: Space Berry

Parent Batch #:; Batch#: 041624-6; Batch Size: g

Sample Received: 04/17/2024; Report Created: 04/23/2024; Expires: 04/23/2025

Manufacturing Date: 04/16/2024

Sampling: ; Environment:

Lic. #00000109ESVM44878444

Harvest Dates: 10/25/2023

Space Dust - Space Berry

Concentrates & Extracts, Diamonds, Extraction Method: CO2

Dispensary License #: ; Manufacturing License #: 00000109ESVM44878444; Cultivation License #: 00000077DCPS00216601





Safety

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

Cannabinoids

85.31%	ND	97.28%
Total THC	Total CBD	Total Cannabinoids Q3

	iotal iiic	IOTAI CDD		Q3	
An	alyte	LOQ	Mass	Mass	Qualifier
		%	%	mg/g	
TH	ICa	0.20	97.28	972.8	D1
Δ9	-THC	0.20	ND	ND	D1
Δ8	-THC	0.20	ND	ND	D1
TH	ICV	0.20	ND	ND	D1
CE	BDa	0.20	ND	ND	D1
CE	BD	0.20	ND	ND	D1
CE	BDV	0.20	ND	ND	D1
CE	BN	0.20	ND	ND	D1
CE	3Ga	0.20	ND	ND	D1
CE	BG .	0.20	ND	ND	D1
CE	SC .	0.20	ND	ND	D1
To	tal		97.28	972.8	

Terpenes

Analyto		100	Mass	Mass	Qualifier
TPL_Terpenes_0	1				

Total THC = THCa * 0.877 + Δ 9-THC Total CBD = CBDa * 0.877 + CBD

LABS

Instrument: HPLC-DAD: ; Method: TPL_Potency_01

Notes:

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

Brian DiMarco **Laboratory Director**

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Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status Q	(ualifier	Analyte	LOQ	Limit	Mass	Status C	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	V1	Hexythiazox	0.48	1.00	ND	Pass	
Acephate	0.19	0.40	ND	Pass	V1	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	L1	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	L1 V1	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	L1	Methiocarb	0.10	0.20	ND	Pass	L1
Carbaryl	0.10	0.20	ND	Pass	L1 V1	Methomyl	0.19	0.40	ND	Pass	L1 V1
Carbofuran	0.10	0.20	ND	Pass	L1 V1	Myclobutanil	0.10	0.20	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass	L1	Naled	0.24	0.50	ND	Pass	L1 V1
Chlorfenapyr	0.48	1.00	ND	Pass		Oxamyl	0.48	1.00	ND	Pass	L1 V1
Chlorpyrifos	0.10	0.20	ND	Pass	L1 V1	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	L1
Cypermethrin	0.48	1.00	ND	Pass		Piperonyl	0.96	2.00	<loq< td=""><td>Pass</td><td></td></loq<>	Pass	
Daminozide	0.48	1.00	ND	Pass		Butoxide	0.70	2.00	LUQ	Pass	
Diazinon	0.10	0.20	ND	Pass	L1 V1	Prallethrin	0.10	0.20	ND	Pass	
Dichlorvos	0.05	0.10	ND	Pass	V1	Propiconazole	0.19	0.40	ND	Pass	M2
Dimethoate	0.10	0.20	ND	Pass	L1 V1	Propoxur	0.10	0.20	ND	Pass	L1 V1
Ethoprophos	0.10	0.20	ND	Pass	L1	Pyrethrins	0.48	1.00	ND	Pass	
Etofenprox	0.19	0.40	ND	Pass	M2	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	V1	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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Heavy Me	tals			Pass	
Analyte	LOQ	Limit	Mass	Status	Qualifier
	PPB	PPB	PPB		
Arsenic	200.0	400.0	ND	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></th></loq<>	Pass	
Mercury	100.0	200.0	<loq< th=""><th>Pass</th><th>L1</th></loq<>	Pass	L1

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

Residu	al Solve	ents				Pass
Analyte		LOQ	Limit	Mass	Status	Qualifier
		PPM	PPM	PPM		
Acetone		460.0	1000.0	ND	Pass	
Acetonitrile		188.6	410.0	ND	Pass	
Benzene		0.9	2.0	ND	Pass	
Butanes		575.0	5000.0	ND	Pass	
Chloroform		27.6	60.0	ND	Pass	
Dichloromet	hane	276.0	600.0	ND	Pass	
Ethanol		2299.9	5000.0	ND	Pass	
Ethyl-Acetat	e	2299.9	5000.0	ND	Pass	
Ethyl-Ether		2299.9	5000.0	ND	Pass	
Heptane		2299.9	5000.0	ND	Pass	
Hexanes		133.4	290.0	ND	Pass	
Isopropyl-Ac	etate	2299.9	5000.0	ND	Pass	
Methanol		1379.9	3000.0	ND	Pass	
Pentanes		133.4	5000.0	ND	Pass	
2-Propanol		2299.9	5000.0	ND	Pass	
Toluene		409.4	890.0	ND	Pass	
Xylenes		92.0	2170.0	ND	Pass	

Performed by GCMS-HS SOP-004. Methods used per AZDHS R9-17-404.03 and the solvent limits set by AZDHS R9-17 Table 3.1. AZDHS approved method for residual solvents by GCMS-HS for all listed analytes. Subcontracted through DVT Registration Certificate Identification Number: 0000031LRCHX78341676

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	

Instruments a DCD/Distings AOAC Matheda 092402, 022202 and 2049, 42

Instrument: qPCR/Plating; AOAC Methods 082102, 022202 and 2018.13 $\,$

Mycotoxins				F	ass
Analyte	LOQ	Limit	Mass	StatusQu	ıalifier
	PPB	PPB	PPB		
B1	8.1	20.0	ND	Pass	M2
B2	8.1	20.0	ND	Pass	M2
G1	8.1	20.0	ND	Pass	M2
G2	8.1	20.0	ND	Pass	M2
Ochratoxin A	8.1	20.0	ND	Pass	M2
Total Aflatoxins	8.1	20.0	ND	Pass	M2



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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.



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