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SROAZ LLC

Sample: 2404TLL0133.0654

Strain: Lazer Fuel

Tempe, AZ 85281

Parent Batch #: 32030624EG-LF; Batch #: 041624-3; 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: 03/06/2024

Space Rocks - Lazer Fuel

Concentrates & Extracts, Caviar, 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

Analyte THCa $\Delta 9$ -THC Δ8-ΤΗС THCV CBDa CBD **CBDV** CBN **CBGa** CBG CBC Total

Forelicy_or		
41.64%	ND	48.30%
Total THC	Total CBD	Total Cannabinoids Q3

48.30%	
otal Cannabinoids	

		•	
LOQ	Mass	Mass	Qualifier
%	%	mg/g	
0.10	46.49	464.9	
0.10	0.87	8.7	
0.10	ND	ND	
0.10	0.82	8.2	
0.10	0.12	1.2	
0.10	ND	ND	
	48.30	483.0	

Terpenes

Analysta	100	Macc	 lace	Qualifier
TPL_Terpenes_01				

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

Instrument: HPLC-DAD: ; Method: TPL_Potency_01

Notes:



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Brian DiMarco Laboratory Director

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Ctatus Oualifar



Pesticides TPL_Pesticides_01

Pass

PPM 0.24 0.19 0.10	PPM 0.50 0.40	PPM ND	Pass			PPM	PPM	PPM		
0.19		ND	Daga					1 1 1 1 1 1		
	0.40		Pass	V1	Hexythiazox	0.48	1.00	ND	Pass	
0.10		ND	Pass	V1	lmazalil	0.10	0.20	ND	Pass	
0.10	0.20	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
0.19	0.40	ND	Pass	L1	Kresoxim	0.10	0.40	ND	Docc	
0.10	0.20	ND	Pass		Methyl	0.17	0.40	ND	F d 5 5	
0.10	0.20	ND	Pass	L1 V1	Malathion	0.10	0.20	ND	Pass	
0.10	0.20	ND	Pass		Metalaxyl	0.10	0.20	ND	Pass	
0.19	0.40	ND	Pass	L1	Methiocarb	0.10	0.20	ND	Pass	L1
0.10	0.20	ND	Pass	L1 V1	Methomyl	0.19	0.40	ND	Pass	L1 V1
0.10	0.20	ND	Pass	L1V1	Myclobutanil	0.10	0.20	ND	Pass	
0.10	0.20	ND	Pass	L1	Naled	0.24	0.50	ND	Pass	L1 V1
0.48	1.00	ND	Pass		Oxamyl	0.48	1.00	ND	Pass	L1 V1
0.10	0.20	ND	Pass	L1 V1	Paclobutrazol	0.19	0.40	ND	Pass	
0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	M2
0.48	1.00	ND	Pass		Phosmet	0.10	0.20	ND	Pass	L1
0.48	1.00	ND	Pass		Piperonyl	0.07	2.00	100	Daga	
0.48	1.00	ND	Pass		Butoxide	0.90	2.00	< LUQ	Pass	
0.10	0.20	ND	Pass	L1 V1	Prallethrin	0.10	0.20	ND	Pass	
0.05	0.10	ND	Pass	V1	Propiconazole	0.19	0.40	ND	Pass	M2
0.10	0.20	ND	Pass	L1 V1	Propoxur	0.10	0.20	ND	Pass	L1 V1
0.10	0.20	ND	Pass	L1	Pyrethrins	0.48	1.00	ND	Pass	
0.19	0.40	ND	Pass	M2	Pyridaben	0.10	0.20	ND	Pass	
0.10	0.20	ND	Pass		Spinosad	0.10	0.20	ND	Pass	
0.10	0.20	ND	Pass	V1	Spiromesifen	0.10	0.20	ND	Pass	
0.19	0.40	ND	Pass		Spirotetramat	0.10	0.20	ND	Pass	
0.19	0.40	ND	Pass		Spiroxamine	0.19	0.40	ND	Pass	
0.48	1.00	ND	Pass		Tebuconazole	0.19	0.40	ND	Pass	
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	
	0.10 0.10 0.10 0.19 0.10 0.10 0.10 0.48 0.10 0.48 0.48 0.48 0.10 0.05 0.10 0.10 0.19 0.10 0.19	0.10	0.10 0.20 ND 0.10 0.20 ND 0.10 0.20 ND 0.10 0.20 ND 0.19 0.40 ND 0.10 0.20 ND 0.48 1.00 ND 0.10 0.20 ND 0.11 0.20 ND 0.10 0.20 ND 0.11 0.20 ND 0.11 0.20 ND 0.12 0.40 ND 0.19 0.40 ND 0.19 0.40 ND 0.48 1.00 ND	0.10 0.20 ND Pass 0.10 0.20 ND Pass 0.10 0.20 ND Pass 0.19 0.40 ND Pass 0.10 0.20 ND Pass 0.48 1.00 ND Pass 0.48 1.00 ND Pass 0.48 1.00 ND Pass 0.10 0.20 ND Pass 0.10 0.20 </td <th>0.10 0.20 ND Pass 0.10 0.20 ND Pass L1 V1 0.10 0.20 ND Pass L1 V1 0.19 0.40 ND Pass L1 0.10 0.20 ND Pass L1 V1 0.10 0.20 ND Pass L1 0.48 1.00 ND Pass L1 V1 0.10 0.20 ND Pass L1 V1 0.10 0.20 ND Pass L1 V1 0.48 1.00 ND Pass L1 V1 0.48 1.00 ND Pass L1 V1 0.05 0.10 ND Pass V1 0.10 0.20 ND Pass L1 0.10</th> <td>0.10 0.20 ND Pass Pass Methyl 0.10 0.20 ND Pass Pass Pass Pass Pass Pass Pass Pass</td> <td>0.10 0.20 ND Pass Pass Methyl 0.19 0.10 0.20 ND Pass Pass Pass Pass Pass Pass Pass Pass</td> <td>0.10 0.20 ND 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V1 Malathion 0.10 0.20 0.10 0.20 ND Pass L1 Methiocarb 0.10 0.20 0.10 0.20 ND Pass L1 V1 Methomyl 0.19 0.40 0.10 0.20 ND Pass L1 V1 Myclobutanil 0.10 0.20 0.10 0.20 ND Pass L1 V1 Myclobutanil 0.10 0.20 0.10 0.20 ND Pass L1 V1 Myclobutanil 0.10 0.20 0.48 1.00 ND Pass L1 V1 Paclobutrazol 0.19 0.40 0.10 0.20 ND Pass L1 V1 Paclobutrazol 0.19 0.40 0.10 0.20 ND Pass Phosmet 0.10 0.20 0.48 1.00 ND Pass </td <td>0.10 0.20 ND Pass pass Methyl 0.19 0.40 ND 0.10 0.20 ND Pass pass L1 V1 Malathion 0.10 0.20 ND 0.10 0.20 ND Pass pass L1 Methiocarb 0.10 0.20 ND 0.10 0.20 ND Pass pass L1 V1 Methiocarb 0.10 0.20 ND 0.10 0.20 ND Pass pass L1 V1 Methiocarb 0.10 0.20 ND 0.10 0.20 ND Pass pass L1 V1 Methiocarb 0.10 0.20 ND 0.10 0.20 ND Pass pass L1 V1 Methiocarb 0.10 0.20 ND 0.48 1.00 ND Pass pass L1 V1 Paclobutrazol 0.19 0.40 ND 0.48 1.00 ND Pass pass Phosmet 0.10 0.20 ND 0.48 1.00 ND</td> <td>0.10 0.20 ND Pass pass Methyl 0.19 0.40 ND Pass pass 0.10 0.20 ND Pass pass pass L1 V1 Malathion pass 0.10 0.20 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0.20 ND Pass pass 0.19 0.40 ND Pass pass L1 Methiocarb pass 0.10 0.20 ND Pass pass 0.10 0.20 ND Pass pass L1 V1 Methomyl pass 0.19 0.40 ND Pass pass 0.10 0.20 ND Pass pass L1 V1 Methomyl pass 0.10 0.20 ND Pass pass 0.10 0.20 ND Pass pass L1 V1 Pass pass 0.24 0.50 ND Pass pass 0.10 0.20 ND Pass pass 0.24 0.50 ND Pass pass 0.10 0.20 ND Pass pass 0.24 0.50 ND Pass pass 0.10 0.20 ND Pass pass Permethrin 0.10 0.20 ND Pass<

 $Instrument: LC\text{-}QQQ \ ; Method: TPL_Pesticides_01$



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Heavy Me	etals				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		468.6	1000.0	ND	Pass	
Acetonitrile		192.1	410.0	ND	Pass	
Benzene		0.9	2.0	ND	Pass	
Butanes		585.8	5000.0	ND	Pass	
Chloroform		28.1	60.0	ND	Pass	
Dichloromet	:hane	281.2	600.0	ND	Pass	
Ethanol		2343.0	5000.0	ND	Pass	
Ethyl-Acetat	:e	2343.0	5000.0	ND	Pass	
Ethyl-Ether		2343.0	5000.0	ND	Pass	
Heptane		2343.0	5000.0	ND	Pass	
Hexanes		135.9	290.0	ND	Pass	
Isopropyl-Ad	cetate	2343.0	5000.0	ND	Pass	
Methanol		1405.8	3000.0	ND	Pass	
Pentanes		135.9	5000.0	ND	Pass	
2-Propanol		2343.0	5000.0	ND	Pass	
Toluene		417.1	890.0	ND	Pass	
Xylenes		93.7	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	

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

Mycotoxins				F	ass
Analyte	LOQ	Limit	Mass	StatusQ	ualifier
	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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