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RSP Productions: Asylum

2985 W. Whitton Ave. Phoenix, AZ 85017 lodellrichardson@gmail.com (702) 338-7088 Lic. #00000070DCBD00783295 Harvest Dates: 04/23/2024

Sample: 2405TLL0154.0770

Strain: Wake N Bake

Parent Batch #:; Batch#: 20240423R8AB1; Batch Size: 15 g

Sample Received: 05/03/2024; Report Created: 05/10/2024; Expires: 05/10/2025

Manufacturing Date:

Sampling: ; Environment:

Wake N Bake Flower

Plant, Flower - Cured

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



Safety

Pass Pesticides

Pass Microbials

Pass

Metals

Cannabinoids

25.31%	
Total THC	

ND	
Total CBD	

	30.07%	
Tota	l Cannabino Q3	ids

Analyte	LOQ	Mass	Mass	Qualifie
	%	%	mg/g	
THCa	0.10	28.01	280.1	
Δ9-ΤΗС	0.10	0.75	7.5	
Δ8-ΤΗС	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	1.16	11.6	
CBG	0.10	0.14	1.4	
CBC	0.10	ND	ND	
Total		30.07	300.7	

Terpenes TPL_Terpenes_01



Instrument: GCMS; Method: TPL_Terp_01





Analyte	LOQ	Mass	Mass	Qualifier
0.14	%	%	mg/g	
β-Myrcene		0.1900	1.900	Q3
β-Caryophyllene		0.1700	1.700	Q3
Ocimene		0.1200	1.200	Q3
α-Humulene		0.1100	1.100	Q3
α-Pinene		0.1100	1.100	Q3
Lina l ool		0.1100	1.100	Q3
β-Pinene		0.1000	1.000	Q3
δ-Limonene		0.0900	0.900	Q3
Terpino l ene		0.0700	0.700	Q3
trans-Nerolidol		0.0400	0.400	Q3
Camphene		0.0100	0.100	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
Total		1.1200	11.200	

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

LABS

Instrument: HPLC-DAD: ; Method: TPL_Potency_01

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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 (Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1L1	Hexythiazox	0.48	1.00	ND	Pass	L1 V1
Abamectin	0.24	0.50	ND	газэ	V1	Imazalil	0.10	0.20	ND	Pass	
Acephate	0.19	0.40	ND	Pass		Imidacloprid	0.19	0.40	ND	Pass	
Acetamiprid	0.10	0.20	ND	Pass		Kresoxim	0.19	0.40	ND	Pass	
Aldicarb	0.19	0.40	ND	Pass		Methy l					
Azoxystrobin	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	
Bifenazate	0.10	0.20	ND	Pass	M1 V1	Metalaxyl	0.10	0.20	ND	Pass	
Bifenthrin	0.10	0.20	ND	Pass	L1 V1	Methiocarb	0.10	0.20	ND	Pass	
Boscalid	0.19	0.40	ND	Pass	V1	Methomyl	0.19	0.40	ND	Pass	
Carbary	0.10	0.20	ND	Pass		Myclobutanil	0.10	0.20	ND	Pass	V1
Carbofuran	0.10	0.20	ND	Pass		Naled	0.24	0.50	ND	Pass	
Chlorantraniliprole	0.10	0.20	ND	Pass		Oxamyl	0.48	1.00	ND	Pass	
Ch l orfenapyr	0.48	1.00	ND	Pass	M2	Paclobutrazol	0.19	0.40	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass		Permethrin	0.10	0.20	ND	Pass	L1 V1
Clofentezine	0.10	0.20	ND	Pass		Phosmet	0.10	0.20	ND	Pass	
Cyfluthrin	0.48	1.00	ND	Pass	M1L1	Piperonyl	0.96	2.00	<loq< td=""><td>Pass</td><td></td></loq<>	Pass	
Cypermethrin	0.48	1.00	ND	Pass	M1L1	Butoxide					
,,	0.40		NID		R1	Prallethrin	0.10	0.20	ND	Pass	M1 V1
Daminozide	0.48	1.00	ND	Pass	M1	Propiconazole	0.19	0.40	ND	Pass	
Diazinon	0.10	0.20	ND	Pass		Propoxur	0.10	0.20	ND	Pass	\/4
Dichlorvos	0.05	0.10	ND ND	Pass		Pyrethrins	0.48 0.10	1.00	<loq ND</loq 	Pass Pass	V1 V1
Dimethoate	0.10	0.20	ND	Pass		Pyridaben	0.10	0.20	ND ND	Pass	٧ı
Ethoprophos	0.10	0.20	ND	Pass Pass	V1	Spinosad Spiromesifen	0.10	0.20	ND	Pass	
Etofenprox Etoxazole	0.19	0.40	ND	Pass	VΙ	Spirotetramat	0.10	0.20	ND	Pass	М1
	0.10	0.20	ND	Pass	M2 V1	Spirotetramat	0.10	0.20	ND	Pass	IVII
Fenoxycarb Fenoyroximate	0.10	0.20	ND	Pass	M ≥ V 1 L1	Tebuconazo l e	0.19	0.40	ND ND	Pass	
Fipronil	0.19	0.40	ND	Pass	LI	Thiacloprid	0.19	0.40	ND	Pass	
Flonicamid	0.19	1.00	ND	Pass	V1	Thiamethoxam	0.10	0.20	ND	Pass	
Fludioxonil	0.46	0.40	ND	Pass	VΙ	Trifloxystrobin	0.10	0.20	ND	Pass	
FIUUIOXOIIII	U. 17	0.40	ND	r'd55		II IIIOXYSUI ODIII	0.10	0.20	ND	r d55	

Instrument: LC-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	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>L1 R1</th></loq<>	Pass	L1 R1

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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- 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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