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Total Health & Wellness dba True Harvest

Sample: 2404TLL0111.0561

Strain: Guava Tangie x Apple Fritz

Parent Batch #: ; Batch#: GTxAFZ0329; Batch Size: 17 g Sample Received: 04/01/2024; Report Created: 04/07/2024; Expires: 04/07/2025

Manufacturing Date:

Sampling: ; Environment:

Lic. #00000100DCWU00857159

jpastor@trueharvestco.com

Harvest Dates:

Phoenix, AZ 85043

Guava Tangie x Apple Fritz

Concentrates & Extracts, Infused/Enhanced Preroll, Extraction Method: Ice/Water Dispensary License #: ; Manufacturing License #: ; Cultivation License #:





Safety

Pass
PesticidesPass
MicrobialsPass
MycotoxinsPass
SolventsPass
MetalsNot Tested
Foreign Matter

Cannabinoids TPL_Potency_01

	38.42%	П	ND		Г	45.05	5%	
	Total THC		Total CBD			Total Canna Q3	abinoids	
A	Analyte		LOQ	Mas	SS	Mass	Qualifie	r
_			%		%	mg/g		
T	HCa		0.10	41.7	'8	417.8		
Δ	∆9-THC		0.10	1.7	8'	17.8		
Δ	\8-THC		0.10	Ν	D	ND		
T	HCV		0.10	Ν	D	ND		
C	BDa		0.10	Ν	D	ND		
C	BD		0.10	Ν	D	ND		
C	CBDV		0.10	N	D	ND		

0.10

0.10

0.10

ND

1.16

0.33

ND

45.05

ND

11.6

3.3

ND

450.5

Terpenes

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A	
—	
Hops	





Analyte	LOQ	Mass	Mass	Qualifier
	%	%	mg/g	
α-Humulene		0.6200	6.200	Q3
β-Caryophyllene		0.6100	6.100	Q3
Linalool		0.3700	3.700	Q3
δ-Limonene		0.2500	2.500	Q3
Guaiol		0.2400	2.400	Q3
trans-Nerolidol		0.2200	2.200	Q3
Ocimene		0.2000	2.000	Q3
β-Myrcene		0.1800	1.800	Q3
β-Pinene		0.1600	1.600	Q3
Terpinolene		0.1500	1.500	Q3
α-Bisabolol		0.1300	1.300	Q3
y-Terpinene		0.1300	1.300	Q3
Eucalyptol		0.0900	0.900	Q3
Caryophyllene Oxide		0.0600	0.600	Q3
α-Pinene		0.0400	0.400	Q3
Camphene		0.0100	0.100	Q3
3-Carene		<	<	Q3
α-Terpinene		<	<	Q3
cis-Nerolidol		<	<	Q3
Geraniol		<	<	Q3
Isopulegol		<	<	Q3
p-Cymene		<	<	Q3

Instrument: GCMS; Method: TPL_Terp_01

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

Instrument: HPLC-DAD: ; Method: TPL_Potency_01

TLABS

CBN

CBG

CBC

Total

CBGa

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Total

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 $\label{linear_property} \textbf{Dispensary License \#: ; Cultivation License \#: } \\$



Pesticides TPL_Pesticides_01

Pass

Analyte	LOQ	Limit	Mass	Status (Qualifier	Analyte	LOQ	Limit	Mass	Status C	Qualifier
	PPM	PPM	PPM				PPM	PPM	PPM		
Abamectin	0.24	0.50	ND	Pass	M1	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	V1	Kresoxim	0.19	0.40	ND	Pass	
Azoxystrobin	0.10	0.20	ND	Pass		Methyl	0.17	0.40	ND	F d 3 3	
Bifenazate	0.10	0.20	ND	Pass		Malathion	0.10	0.20	ND	Pass	L1 V1
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	V1
Chlorfenapyr	0.48	1.00	ND	Pass	M2	Oxamyl	0.48	1.00	ND	Pass	
Chlorpyrifos	0.10	0.20	ND	Pass	V1 M2	Paclobutrazol	0.19	0.40	ND	Pass	
Clofentezine	0.10	0.20	ND	Pass	V1	Permethrin	0.10	0.20	ND	Pass	M2
Cyfluthrin	0.48	1.00	ND	Pass	M1	Phosmet	0.10	0.20	ND	Pass	11
Cypermethrin	0.48	1.00	ND	Pass		Piperonyl	0.96	2.00	ND	Pass	
Daminozide	0.48	1.00	ND	Pass		Butoxide	0.70		ND		
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	V1
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	V1
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	M1
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	M2 I1

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



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Heavy Me	tais				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	

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		498.5	1000.0	ND	Pass	
Acetonitrile		204.4	410.0	ND	Pass	
Benzene		1.0	2.0	ND	Pass	
Butanes		623.1	5000.0	ND	Pass	
Chloroform		29.9	60.0	ND	Pass	
Dichloromet	hane	299.1	600.0	ND	Pass	
Ethanol		2492.5	5000.0	ND	Pass	
Ethyl-Acetat	e	2492.5	5000.0	ND	Pass	
Ethyl-Ether		2492.5	5000.0	ND	Pass	
Heptane		2492.5	5000.0	ND	Pass	
Hexanes		144.6	290.0	ND	Pass	
Isopropyl-Ad	etate	2492.5	5000.0	ND	Pass	
Methanol		1495.5	3000.0	ND	Pass	
Pentanes		144.6	5000.0	ND	Pass	
2-Propanol		2492.5	5000.0	ND	Pass	
Toluene		443.7	890.0	ND	Pass	
Xylenes		99.7	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	Pass
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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