

VT Cheddar

 Sample ID: BIA251010S0255
 Strain: VT Cheddar

 Produced:
 Collected:
 Received: 10/10/2025
 Completed: 10/17/2025
 Batch#: SCLT0161-19 F

 Client
King Cola
 Lic. # SCLT0161
 Po Box 17
 Williamstown, VT 05679

 Matrix: Plant
 Type: Flower - Cured
 Sample Size: 8.05 g
 Lot#:


Summary

Test	Date Tested	Result
Sample		Complete
Cannabinoids	10/14/2025	Complete
Moisture	10/13/2025	11.10% - Complete
Water Activity	10/13/2025	0.574 aw - Complete
Terpenes	10/14/2025	Complete
Microbials	10/17/2025	Complete

Cannabinoids

Completed

24.81% Total THC					0.09% Total CBD			30.57% Total Cannabinoids			
Analyte	LOQ	Results	Results	Mass	Analyte	LOQ	Results	Results	Mass		
	mg/g	%	mg/g	mg/serving		mg/g	%	mg/g	mg/serving		
CBDVa	0.0003	<LOQ	<LOQ		CBCVa	0.0003	<LOQ	<LOQ			
CBDV	0.0003	<LOQ	<LOQ		CBNa	0.0003	0.06	0.6			
CBDa	0.0005	0.10	1.0		Δ9-THC	0.0005	1.61	16.1			
CBGa	0.0005	0.95	9.5		Δ8-THC	0.0003	0.04	0.4			
CBG	0.0005	0.11	1.1		Δ10-THC*	0.0002	<LOQ	<LOQ			
CBD	0.0005	<LOQ	<LOQ		CBL	0.0005	<LOQ	<LOQ			
THCV	0.0003	0.07	0.7		CBC	0.0003	0.30	3.0			
CBLV	0.0003	0.06	0.6		THCa	0.0005	26.45	264.5			
CBCV	0.0003	<LOQ	<LOQ		CBCa	0.0006	0.62	6.2			
THCVa	0.0003	0.21	2.1		CBLa	0.0005	<LOQ	<LOQ			
CBN	0.0005	<LOQ	<LOQ		Total THC		24.81	248.05			
					Total CBD		0.09	0.85			
					Total		30.57	305.74	0.00		

Analyst: 056

Cannabinoids Methodology: High Performance Liquid Chromatography (HPLC) using PerkinElmer FLEXAR™ with Photo Diode Array Detector (PDA)

Total CBD and total THC are calculated values, to account for assumed decarboxylation from the acid form (THCA or CBDA) to the neutral form, causing weight loss of the acid group. These values are calculated as follows:

$$\text{Total THC} = (\text{THCA} \times 0.877) + \Delta 9\text{-THC}$$

$$\text{Total CBD} = (\text{CBDA} \times 0.877) + \text{CBD Reagent}$$

Blanks: < LOQs for all analytes

LOQ = The lowest quantity that this method can reliably detect. Any cannabinoid that was not detected is assumed to be less than the stated LOQ (<LOQ).

All results reflect dry weight of material, based on % moisture of the sample.

Measurement of Uncertainty (MU): the parameter, associated with the result of a measurement, that characterizes the dispersion of the values that could reasonably be attributed to the

particular quantity subject to measurement. Δ9-THC MU = ±0.005% Total THC MU = ±0.007%

All other cannabinoid MU values are available upon request.

All moisture and water activity analysis is determined by dewpoint measurement using an AQUALAB water activity meter.

*The result is the sum of delta-10 isomers.




 Luke Emerson-Mason
 Laboratory Director
 10/17/2025

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Terpenes

Completed

Analyte	LOQ	Results	Results
	mg/g	mg/g	%
Limonene	0.010	3.265	0.327
Ocimene	0.010	2.977	0.298
β-Myrcene	0.010	2.626	0.263
Linalool	0.010	2.283	0.228
β-Caryophyllene	0.010	2.223	0.222
β-Pinene	0.010	1.584	0.158
α-Pinene	0.010	1.082	0.108
α-Humulene	0.010	0.924	0.092
Eucalyptol	0.010	0.245	0.025
Camphene	0.010	0.177	0.018
Terpinolene	0.010	0.143	0.014
γ-Terpinene	0.010	0.055	0.006
α-Terpinene	0.010	0.030	0.003
3-Carene	0.010	<LOQ	<LOQ
α-Bisabolol	0.010	<LOQ	<LOQ
Caryophyllene Oxide	0.010	<LOQ	<LOQ
cis-Nerolidol	0.010	<LOQ	<LOQ
Geraniol	0.010	<LOQ	<LOQ
Guaiol	0.010	<LOQ	<LOQ
Isopulegol	0.010	<LOQ	<LOQ
p-Cymene	0.010	<LOQ	<LOQ
trans-Nerolidol	0.010	<LOQ	<LOQ
Total		17.614	1.761

Primary Aromas



Analyst: 048

LOQ = The lowest quantity this method can reliably detect. Any terpene that was not detected is assumed to be less than the stated LOQ (<LOQ).

Terpene Methodology: Headspace Sampler, Gas Chromatography-Mass Spectrometry (GC-MS), using Perkin Elmer Clarus® SQ8 GC MS

Reagent Blanks: < LOQs for all analytes

All results reflect dry weight of material, based on % moisture of the sample.

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Pathogens

Completed

Pathogens	LOD CFU/g	Results CFU/g
Aspergillus	5	Not Detected
Shiga Toxin E. Coli	5	Not Detected
Salmonella SPP	5	Not Detected

Analyst: 049

Test Methodology: Bio-Rad IQ-Check PCR Kits

cfu/g = colony forming units per gram

LOD = The lowest quantity that this method can reliably detect. Any microbial growth that was not detected is assumed to be less than the stated LOD (<LOD).

Reagent Blanks: <LOD for all analytes




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