



The Good Lab

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Certificate of Analysis

Customer ID	750	Cust Name	Hedberg Fence		
Sample ID	1902766B	Sample Name	Flower		
Sample Type	Biomass	Date Received	11/15/2019	Date Completed	11/22/2019

Total THC %	Total CBD %	Total Cannabinoids %	Potential Δ9THC %	Potential CBD %
0.27	12.52	13.29	0.27	11.01

Sample Preparation Method to determine Δ9-THC: CDA Decarb

Potency Profile Percentage		Terpene Profile Percentage			
CBDV	0.00	Alpha-Pinene	NOT TESTED	Terpinolene	NOT TESTED
CBDA	12.30	Camphene	NOT TESTED	Linalool	NOT TESTED
CBGA	0.42	Beta-Pinene	NOT TESTED	(-)-Isopulegol	NOT TESTED
CBG	0.00	Beta-Myrcene	NOT TESTED	Geraniol	NOT TESTED
THCV	0.00	Delta-3-Carene	NOT TESTED	Beta-Caryophyllene	NOT TESTED
CBD	0.22	Alpha-Terpinene	NOT TESTED	Alpha-Humulene	NOT TESTED
CBN	0.00	Limonene	NOT TESTED	cis-Nerolidol	NOT TESTED
Δ9THC	0.27	Alpha-Ocimene	NOT TESTED	trans-Nerolidol	NOT TESTED
CBC	0.08	Eucalyptol	NOT TESTED	(-)-Guaiaol	NOT TESTED
THCA	0.00	Beta-Ocimene	NOT TESTED	(-)-Caryophyllene Oxide	NOT TESTED
TOTAL Cannabinoids	13.29	Gamma-Terpinene	NOT TESTED	Alpha-Bisabolol	NOT TESTED
Total THC = Δ9THC + THCA + THCV Total CBD = CBD + CBDA + CBDV Potential Δ9THC = Δ9THC + (THCA x .877) Potential CBD = CBD + (CBDA x .877) LOQ = Limit of Quantitation TR = Trace ND = Not Detected This product has been analyzed by The Good Lab using valid testing methodologies developed by Colorado state certified cannabis laboratories.				TOTAL Terpenes	0.0000
		Heavy Metals		NOT TESTED	
		Residual Solvents		NOT TESTED	
		Mycotoxins		NOT TESTED	
		Pesticides		NOT TESTED	

FINAL APPROVAL

Analysis:

Gregory P. Duran, Lab Owner

Quality Control:

M. Teri Robnett, Lab Manager

Thank you for choosing **The Good Lab** for your analytical needs. This report outlines the results of your product analysis. If you have any further questions regarding your product, feel free to contact us for a consultation at (720) 245-8323 or goodlabcolorado@gmail.com.

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Potency test results are reported in percentage by **dry weight**. Detectable amounts below .06% are shown as TR (trace) or <LOQ. Our standard detection limit is .02%. Results below .02% are considered unreliable and are reported as zero (0.00) or Not Detected (ND).