Analysis of contaminants in hemp using LC and GC coupled to MS/MS

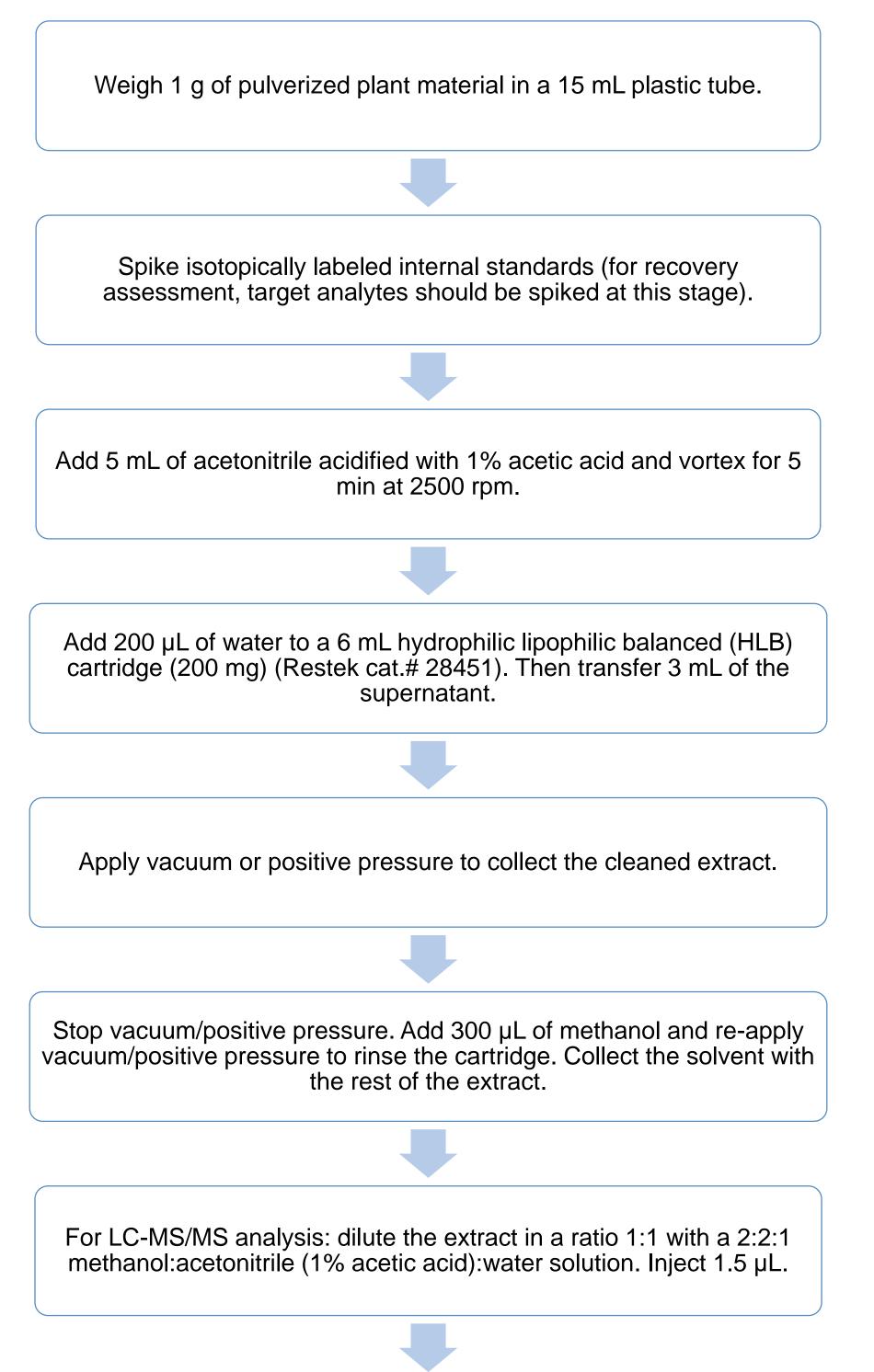
Introduction

- Hemp is a class of Cannabis sativa that contains significantly lower levels of tetrahydrocannabinol (THC), and may have higher levels of cannabidiol and cannabigerol (CBD and CBG).
- Like cannabis and other crops, dried hemp plant material may contain various contaminants that are harmful to humans.
- The high complexity of hemp and cannabis samples, and the broad range of contaminants being regulated at minimum required performance levels (MRPL) in the order of parts per billion (ppb), demands for robust, fast, and effective analytical methods.
- This work describes a complete workflow for the analysis of diverse contaminants in hemp using hydrophilic lipophilic balanced (HLB) cartridges to clean-up organic hemp extracts, and using LC and GC coupled to MS/MS for reliable instrumental analysis.

Goal

To provide an effective workflow for the analysis of pesticides and mycotoxins in hemp and cannabis plant material by using a single extract and LC-MS/MS and GC-MS/MS.

Method development: sample preparation



For GC-MS/MS analysis: transfer 1 mL of cleaned supernatant to a dSPE tube containing magnesium sulfate and C18 (cat.# 26242). Vortex briefly and centrifuge for 5 min. Dilute the extract in a ratio 1:1 with a 1:1 hexane:acetone (1% acetic acid) solution. Inject 1µL.

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Method development: LC/GC-MS/MS

Table 1. LC-MS/MS conditions (ionization: ESI)

Table 1. LC-WS/WS COnditions (IONZATION. ESI)								
Column	Raptor ARC-18 2.7 µm, 150 mm x 2.1 mm (cat.# 9314A62)							
Guard Column	Raptor ARC-18 EXP Guard Column Cartridge 2.7 µm, 5 x 2.1 mm (cat.# 9314A0252)							
Mobile Phase A	Water, 2 mM ammonium formate, 0.1% formic acid							
Mobile Phase B	Methanol, 2 mM ammonium formate, 0.1% formic acid							
Time Program	<u>Time (min.)</u>	<u>%B</u>	<u>Time (min.)</u>	<u>%B</u>				
	0	5	11	75				
	1.0	50	11.5	80				
	2.5	50	13.5	80				
	4.0	65	15.5	95				
	7.0	65	16.5	100				
	7.5	70	19.5	100				
	9.0	70	19.6	5				
	9.5	75						
Other	Column T: 40°C; autosampler T: 10°C; flow: 0.4							
parameters	mL/min; injection volume: 1.5 µL							
Instrument	Shimadzu LCMS-8045							

Table 2. GC-MS/MS conditions (ionization: EI)

GC ColumnRxi-5ms 30 m x 0.25 mm x 0.25 µm (cat.# 13423)InjectionSplitless, 1 µL (0.5 min splitless time, 7 mL/min split flow)LinerTopaz 4.0 mm ID Single Taper Inlet Liner w/ Wool (cat.# 23447)Inj. T250°CPurge Flow5 mL/minOven70°C (hold 1 min) to 220°C by 30°C/min; to 240°C by 5°C/min; to 315°C (hold 10 min) by 10°C/minCarrier GasHe, at a constant flow of 1.4 mL/min
Injectionflow)LinerTopaz 4.0 mm ID Single Taper Inlet Liner w/ Wool (cat.# 23447)Inj. T250°CPurge Flow5 mL/minOven70°C (hold 1 min) to 220°C by 30°C/min; to 240°C by 5°C/min; to 315°C (hold 10 min) by 10°C/min
Liner(cat.# 23447)Inj. T250°CPurge Flow5 mL/minOven70°C (hold 1 min) to 220°C by 30°C/min; to 240°C by 5°C/min; to 315°C (hold 10 min) by 10°C/min
Purge Flow5 mL/minOven70°C (hold 1 min) to 220°C by 30°C/min; to 240°C by 5°C/min; to 315°C (hold 10 min) by 10°C/min
Oven70°C (hold 1 min) to 220°C by 30°C/min; to 240°C by 5°C/min; to 315°C (hold 10 min) by 10°C/min
5°C/min; to 315°C (hold 10 min) by 10°C/min
Carrier Gas He, at a constant flow of 1.4 mL/min
Transfer line T 290°C
Source T 330°C
Instrument Thermo Trace 1310-TSQ 8000

Results and discussion

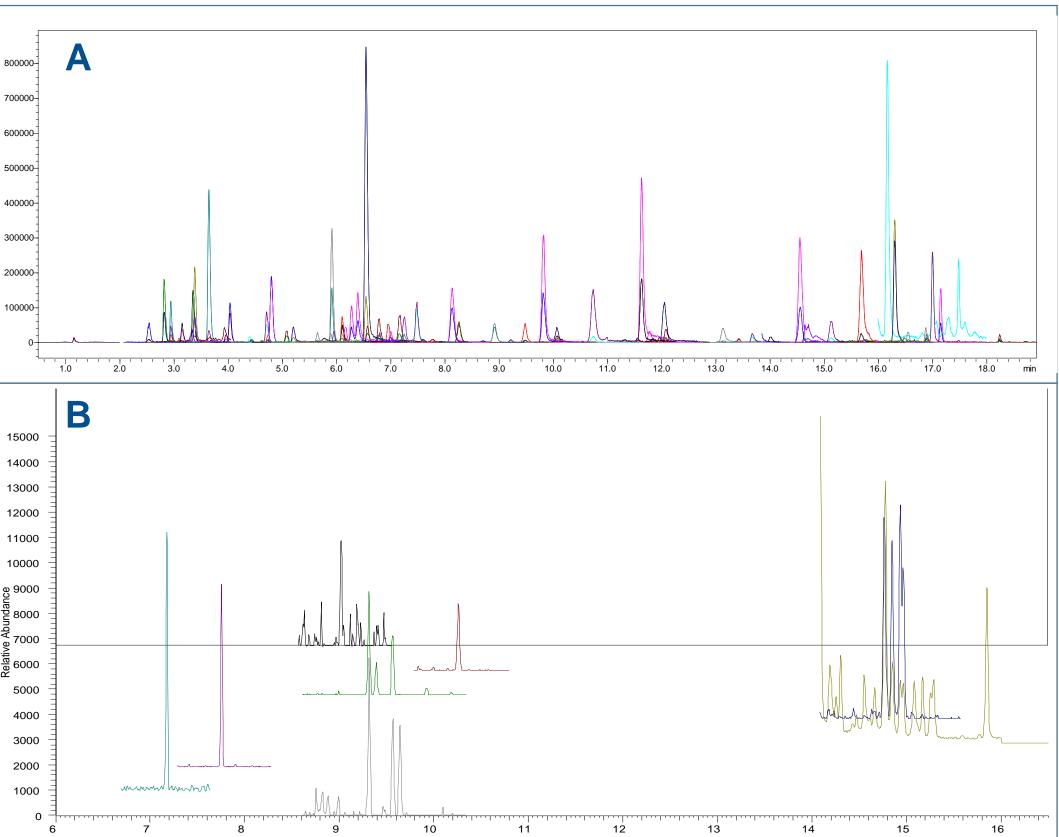


Figure 2. Chromatograms corresponding to LC (A) and GC (B) amenable contaminants extracted from a hemp sample spiked at 100 ng/g

Table 3. Figures of merit corresponding to pesticides and mycotoxins analyzed in hemp (CBG variety) 0.1 µg/g Canada California Method (n=4) Action R^2 Action LOQ, Contaminant Con level, Accuracy µg/g level, µg/g (RSD) µg/g Daminozide 0.02 0.9974 80 (6) 0.1 0.1 Spinosad-0.005 0.9996 0.02 100 (4) Diazinon 0.1 0.005 0.9994 Thiamethoxam 0.02 102 (3) 5 Coumapho 0.005 0.9989 105 (5) Clofentezir 0.05 0.005 96 (2) 0.5 0.9980 Spinosad 3 0.01 0.02 0.9983 93 (11) Spinetoran Imidacloprid 5 0.1 0.02 0.005 0.9979 101 (2) Spinetoran Dimethoate 0.1 0.005 0.9993 100 (4) Trifloxystro Acetamiprid 0.1 Prallethrin Thiacloprid 0.02 0.005 0.9993 100 (8) 0.1 0.005 0.9977 101 (5) Hexythiazo 0.1 0.02 0.9988 83 (12) 0.1 Cyfluthrin 0.1 0.02 0.9976 99 (5) Mevinphos (I, II) 0.1 0.05 Etoxazole 0.005 0.9983 102 (5) Chlorpyrifo Carbofuran 0.1 0.02 0.02 0.9979 96 (6) 0.5 0.05 Permethrir 0.9983 99 (12) Dichlorvos 0.1 0.1 0.1 Fenpyroxir 0.005 0.02 0.9982 100 (3) 0.1 Bifenthrin 0.005 Chlorantraniliprole 0.02 0.9953 94 (3) 10 Abamectin 0.1 0.05 0.01 0.9987 80 (11) Cypermeth 0.005 0.9992 105 (5) Etofenprox 0.02 0.005 0.9967 0.02 105 (4) Pyridaben 0.1 Azoxystrobin 0.005 0.9959 Myclobutani 0.1 0.02 100 (4) Acequinoc 0.005 0.9991 79 (16) 0.02 Flonicamic 0.1 Spiroxamine 0.005 0.9920 72 (15) Fipronil 0.1 0.1 0.005 0.02 0.9988 89 (5) Fludioxoni Fenoxycarb 0.1 Aflatoxin (0.005 0.9958 97 (3) Methiocarb 0.1 0.02 0.02 0.9981 Aflatoxin (Spiromesifen 0.1 108 (12) Aflatoxin E 0.02 0.9949 86 (8) 0.02 0.1 0.02 0.01 0.9957 Aflatoxin E 0.1 102 (7) Paclobutrazol 0.01 Ochratoxir 0.02 0.9954 97 (4) 0.5 0.01 Captan (G 0.05 0.9955 93 (8) Dimethomorph (I,II) 2 Chlordane 0.005 0.9988 0.05 0.1 101 (4) Tebuconazole Chlorfenap 0.005 0.1 0.9970 100 (5) Bifenazate 0.02 0.02 0.9942 79 (12) Methyl par 0.1 Fenhexamid 0.02 0.9985 PCNB (GC 98 (18) 0.1 0.1 Propiconazole 0.005 0.9993 Cyfluthrin 0.02 98 (4) 0.1 Spirotetramat 0.005 0.1 0.02 0.9995 Ethoprophos 98 (5) Cypermetl 0.02 0.02 0.9980 0.1 91 (8) Kresoxym-methyl

Acephate Methomy Oxamyl Aldicarb Naled Carbaryl Propoxur Imazalil Metalaxyl Phosmet Boscalid Malathion

• Hemp and cannabis extracts are characterized for having a high concentration of hydrophobic constituents. By mixing 3 mL of extract with 200 µL of water prior to SPE clean-up with HLB it was possible to remove major hydrophobic interferences. The addition of 300 μ L of methanol helped in the elution of all target pesticides.

• Calibration curves to cover a range of 0.005 and 1.5 µg/g in matrix (10 points) were prepared by post-spiking blank hemp extract with target analytes at various concentrations, and internal standards (9 compounds). All analytes, except clofentezine and captan, showed $R^2 > 0.99$. It is recommended to use deuterated analogues for these two compounds.

• Accuracy and precision were assessed by spiking hemp samples at 0.01, 0.05, 0.1, and 0.5 μ g/g (n=4), and estimating their concentration using the calibration curve prepared in hemp extract. Accuracy and precision values for the great majority of pesticides were within 70 - 130% and below 30%, respectively.

• Sample prep, extract dilution, injection volume, chromatographic separation were all critical in resolving analytes from interferences as to minimize possible matrix effects and reach the required MRPLs.

• In total 9 deuterated analytes were used to account for sample prep and instrumental variation.

• The use of dSPE containing magnesium sulfate was essential to remove any water left in extracts after the first clean-up step.

• Pyrethrins I and II, and piperonyl butoxide were present in the hemp samples used for this work, so they were excluded from the table.

• Overall, the proposed workflow showed satisfactory results in the quantification of all target pesticides and mycotoxins. For the great majority of the compounds the LOQ values were significantly below the action levels established by the state of CA in inhalable cannabis, and comply with Canada regulations.

An easy and effective workflow for the analysis of pesticides and mycotoxins in hemp was developed. Satisfactory results in terms of figures of merit (LOQs, R², accuracy, and precision) were obtained for the great majority of target contaminants.

• Reyes-Garcés N, Myers C. J Sep Sci. 2021; 44: 2564-2576. https://doi.org/10.1002/jssc.202001265

ntaminant	California Action level, µg/g	Canada Action level,	Method LOQ, µg/g	R^2	0.1 µg/g (n=4) Accuracy			
		μg/g			(RSD)			
spinosyn A	0.1*	0.1*	0.00355	0.9963	74 (14)			
opinooyn	0.1	0.02	0.005	0.9990	96 (4)			
OS	0.1	0.02	0.01	0.9989	84 (9)			
ne	0.1	0.02	0.02	0.9844	38 (12)			
- spinosyn D	0.1*	0.1*	0.0029	0.9961	74 (20)			
m - spinosyn J	0.1^	0.02^	0.0042	0.9960	73 (16)			
m - spinosyn L	0.1^	0.02^	0.001	0.9959	75 (13)			
obin	0.1	0.02	0.005	0.9987	106 (6)			
<u> </u>	0.1	0.05	0.05	0.9993	98 (9)			
ОХ	0.1	0.01	0.005	0.9915	75 (26)			
	2	0.2	0.15	0.9943	-			
	0.1	0.02	0.005	0.9965	90 (6)			
OS	0.1	0.04	0.02	0.9961	97 (7)			
ns	0.5	0.5	0.01	0.9979	77 (6)			
mate	0.1	0.02	0.005	0.9977	97 (7)			
	3	1	0.005	0.9974	93 (7)			
nB1a	0.1	0.1	0.01	0.9973	94 (11)			
hrin	1	0.3	0.075	0.9966	90 (10)			
x	0.1	0.05	0.005	0.9977	78 (6)			
l	0.1	0.05	0.005	0.9991	92 (7)			
cyl	0.1	0.03	0.02	0.9961	80 (16)			
d	0.1	0.05	0.05	0.9968	87 (17)			
	0.1	0.06	0.01	0.9985	106 (10)			
il	0.1	0.02	0.02	0.9958	83 (6)			
G2	0.02#	0.002	0.02	0.9905	87 (6)			
G1	0.02#	0.002	0.005	0.9969	88 (12)			
32	0.02#	0.002	0.01	0.9958	87 (8)			
31	0.02#	0.002	0.005	0.9959	85 (7)			
n A	0.02	0.02	0.02	0.9966	83 (8)			
SC)	0.7	_	0.075	0.9829	78 (25)			
e (GC)	0.1	_	0.02	0.9941	81 (1)			
pyr (GC)	0.1	0.05	0.02	0.9939	98 (9)			
rathion (GC)	0.1	0.05	0.005	0.9969	97 (2)			
C)	0.1	0.02	0.01	0.9965	81 (8)			
(GC)	2	0.2	0.02	0.9927	92 (6)			
hrin (GC)	1	0.3	0.05	0.9897	97 (7)			
tal spinosad: $\Delta MPPL$ for total spinoteram: # $\Lambda G2 + \Lambda G1 + \Lambda B1 + \Lambda B2 < 0.002 \mu a/a$								

*MRPL for total spinosad; ^MRPL for total spinoteram; # AG2+AG1+AB1+AB2<0.002µg/g

Conclusions

References