### Chemical Comprehensive Analysis of *Ephedra sinica* Extract, Lot No. RK-3-28-1-ES for the Botanical Safety Consortium

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#### Submitted by:

#### Lori L. Smith and Kristin L. Aillon

MRIGlobal 425 Dr Martin Luther King Jr Boulevard Kansas City, MO 64110-2241

#### Chemical Information: Ephedra sinica Extract

CAS No.: 85940-38-1	Lot No.: RK-3-28-1-ES
DTXSID No.: DTXSID801018482	Amount Received: 1 × ~ 153000 mg
	Sample Receipt Date: 10/7/21
	Appearance: Black Solid
	Supplier: China Medical University (Aerial parts, ground) Post-Handling Supplier: University of Mississippi (95% Ethanol extract) Receipt Condition: Ambient, packaging intact Shipping Containers: Clear-glass jar Storage Condition at MRIGlobal: - 20°C under inert gas

#### **Reference Standards**

Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure: Ephedrine hydrochloride (as (L)-Ephedrine) 50-98-6 USP (Rockville, MD) R08630 99.9% per C of A  $C_{10}H_{15}NO \cdot HC1$ 201.69 HCl HCl $CH_3$ 

Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure: Pseudoephedrine hydrochloride (as Pseudoephedrine) 345-78-8 USP (Rockville, MD) R110F0 100.0% per C of A C<sub>10</sub>H<sub>15</sub>NO·HCl 201.69



Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure: DL-Norephedrine hydrochloride (as Norephedrine) 154-41-6 Sigma-Aldrich (St. Louis, MO) 115K0718 >99% per C of A C9H13NO·HCl 187.67 OH HCl



Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure: Norpseudoephedrine 492-39-7 Cayman Chemical (Ann Arbor, MI) 0511220 99.4% per C of A C<sub>9</sub>H<sub>13</sub>NO 151.21



Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure: N-Methylephedrine 552-79-4 Sigma-Aldrich (St. Louis, MO) BCBL4243V 99.6% per C of A C<sub>11</sub>H<sub>17</sub>NO 179.26



Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure:

(D)-Catechin 154-23-4 USP (Rockville, MD) R13070 97% per C of A C<sub>15</sub>H<sub>14</sub>O<sub>6</sub> 290.27



Name: CAS No.: Supplier: Lot No.: Purity: Molecular Formula: Molecular Weight: Structure:	Quinoline 91-22-5 Sigma-Aldrich (St. Louis BCBX3571 98.7% per C of A C <sub>9</sub> H <sub>7</sub> N 129.16	e, MO)
Prepare <i>Ephedra sinica</i> extract solutions in triplicate		
<ol> <li>Vortex-mix Ephedra in 95% ethanol (2.5</li> <li>Sonicate 20 min.</li> <li>Shake 30 min.</li> <li>Add deionized wate</li> <li>Centrifuge (500×g, 4</li> <li>Filter (0.22-um Nylog)</li> </ol>	a <i>sinica</i> extract (~ 10 mg) mL). er (2.5 mL) and mix. 5 min).	
	>	Identification and CAD Estimation
Prepare 3 spiked Ephedra sinica extract solutions sets in triplicate		
Extract (	e and vortex. (450 μL)► solution (50 μL)	Standard Addition Quantitation Norephedrine, Quinoline*
Set B <u>Combine and v</u>	ortex:	*Qualitative evaluation in unspiked solution
10x Diluted Ext Spiking solution 50% aq. ethance	rract (10 μL) n (50 μL)► bl (440 μL)	Standard Addition Quantitation (L)-ephedrine and Pseudoephedrine
Set A <u>Combine and vortex</u> : Extract (10 µL) Spiking solution (50 µl 50% aq. ethanol (440	L)▶	Standard Addition Quantitation Norpseudoephedrine, N-Methylephedrine, (D)-Catechin

Figure 1. Ephedra sinica extract solution preparation flowchart.

Instrumentation:						
Liquid Chromatograph:	Acquity UPLC with additional post-column inverse gradient Acquity UPLC pump (Waters Corporation; Milford, MA)					
Photodiode Array Detector (UV):	Acquity PDA Detector (Waters Corporation; Milford, MA)					
Charged Aerosol Detector (CAD):	Corona Veo RS (Thermo Fisher Scientific; Waltham, MA)					
High Resolution Mass Spectrometer (HRMS):	Xevo G2-XS (Waters Corporation; Milford, MA)					

# Table 1. UPLC-UV-CAD/HRMS System

### Table 2. Identification and CAD Estimation - Acquisition Parameters

UPLC Parameters:	
Caluman	Avantor ACE Excel C18-PFP, 1.7 µm, 100x 2.1 mm (Advanced
Column:	Chromatography Technologies, UK)
Autosampler Temperature:	6°C
Column Temperature:	30°C
Injection Volume:	10 µL
Flow Rate:	0.3 mL/min
Mobile Phase A:	1% (v/v) Acetic Acid in Water
Mobile Phase B:	1% (v/v) Acetic Acid in Acetonitrile
	Forward: 0% B held for 0.5 min; ramp to 10% B over 4 min; ramp to 55%
	B over 45 min; ramp to 100% B over 18 min and hold for 2.5 min;
	decreased to 0% B in 1 min and held for 4 min.
Gradient Elution Profiles:	
	Inverse: 100% B held for 0.5 min; decrease to 90% B over 4 min;
	held for 2.5 min; ramped to 100% B in 1 min and held for 4 min.
Run Time:	75 minutes
UV (PDA) Parameters:	
2D Channel Wavelength:	254 nm
3D Channel Wavelength Range:	190 to 400 nm, 1.2 nm resolution
CAD Parameters:	· · · · ·
Sampling Rate:	10 points/sec
Scale Factor:	1000
Evaporation Temperature:	35°C
Scale/Power Function Value:	20 pA/1.0
Nitrogen	57.5 psi
HRMS Parameters:	
Ionization/Acquisition Mode:	ESI+; MSe sensitivity mode
Acquisition Range:	<i>m</i> /z 100 to <i>m</i> /z 1200
Scan Time:	0.3 s
Lock Mass:	Leucine encephalin
	(m/z 278.1135, 556.2766; Scan for 0.200 s, interval: 45 s)
Capillary Voltage:	2.5 kV
Cone Voltage:	40 V
Collision Energy:	Low CE: 6 eV High CE: 20-30 eV
Source Temperature:	110°C
Desolvation Temperature:	250°C
Cone Gas Flow:	50 L/h
Desolvation Gas Flow:	600 L/h
Data Acquisition System:	UNIFI version 1.9.4

## Identification



**Figure 2.** Partial chromatograms from the UPLC-UV-CAD/HRMS analysis of *Ephedra sinica* extract showing the majority of analyte peaks from the 75-min analysis. Peak numbers correlate with results presented in Table 3.

Peak Number	Retention Time (min)	Area (%)ª	Observed Neutral Mass <sup>b</sup> (Da)	Mass Error (mDa)	Component ID	CAS	Molecular Formula	Structure	Confidence Level <sup>c-h</sup>	Additional Comments
1	5.31	10.29	151.0994	-0.3	Norephedrine	14838-15-4	C <sub>9</sub> H <sub>13</sub> NO	OH CH <sub>3</sub> CH <sub>3</sub> NH <sub>2</sub>	1	Standard Addition Quantitation (see Table 12).
2	5.65	9.38	151.0997	0.0	Norpseudoephedrine	492-39-7	C <sub>9</sub> H <sub>13</sub> NO	OH CH <sub>3</sub> CH <sub>3</sub> NH <sub>2</sub>	1	Standard Addition Quantitation (see Table 12).
3	5.89	1.28	313.1536	-14.2	Armepavine	524-20-9	C <sub>19</sub> H <sub>23</sub> NO <sub>3</sub>	OH O O O O O O O H 3	3	CAD Estimation (see Table 4).
4	6.28	37.88	165.1155	0.1	(L)-Ephedrine	299-42-3	C <sub>10</sub> H <sub>15</sub> NO	CH <sub>3</sub>	1	Standard Addition Quantitation (see Table 12).
5	6.73	16.21	165.1157	0.4	Pseudoephedrine	90-82-4	C <sub>10</sub> H <sub>15</sub> NO	CH <sub>3</sub>	1	Standard Addition Quantitation (see Table 12).
6	7.35	8.19	179.1315	0.5	N-methylephedrine	552-79-4	C <sub>11</sub> H <sub>17</sub> NO	OH CH <sub>3</sub> NCH <sub>3</sub> CH <sub>3</sub>	1	N-methylephedrine Standard Addition Quantitation (see Table 12). Possible coelution with Pseudomethyl- ephedrine.

Table 3. Identification Summary for *Ephedra sinica* Extract Components

7 <sup>i</sup>	8.12	1.16	290.0798	0.8	(D)-Catechin	154-23-4	C <sub>15</sub> H <sub>14</sub> O <sub>6</sub>	HO OH OH	1	(D)-Catechin Standard Addition Quantitation (see Table 12). Possible coelution with Epicatechin.
8	8.28	1.28	<i>m/</i> z 265.1553	NA	Unknown	NA	NA	NA	5	Singly charged ion observed based on isotope distribution
9	9.39	3.62	<i>m/</i> z 220.0611	0.63	Unknown	NA	$C_{11}H_9NO_4$	NA	4	NA
10A <sup>j</sup>	11.89	1.33	274.0813	-2.8	Afzelechin and/or Epiafzelechin	2545-00-8 and/or 36801-69-1	C <sub>15</sub> H <sub>14</sub> O <sub>5</sub>		3	CAD Estimation (see Table 4); coelutes with Peak 10B.
10B <sup>j</sup>	11.99	1.33	<i>m/</i> z 565.1589	NA	Unknown	NA	NA	NA	5	Singly charged ion observed based on isotope distribution; coelutes with Peak 10A.
11	13.63	3.70	416.1004 and 578.1644	-10.3 and 0.8	Apigenin- 5-rhamnoside and Vitexin rhamnoside	Unassigned and 32426-34-9	C <sub>21</sub> H <sub>20</sub> O <sub>9</sub> and C <sub>27</sub> H <sub>30</sub> O <sub>14</sub>		3	CAD Estimation (see Table 4). Peak 11 has a leading shoulder indicating the presence of coeluting compounds.

12	13.81	0.35	432.1067	1.1	Kaempferol-3-O- rhamnoside	Unassigned	C <sub>21</sub> H <sub>20</sub> O <sub>10</sub>		3	CAD Estimation (see Table 4).
13	16.45	1.11	302.0442	1.6	Herbacetin and/or Quercetin	527-95-7 and/or 117-39-5	C <sub>15</sub> H <sub>10</sub> O7	HO OH OH OH OH O OH HO OH OH HO OH OH OH O	3	CAD Estimation (see Table 4).
14	17.00	0.58	<i>m/z</i> 679.5142 and <i>m/z</i> 340.2611	NA	Unknown	NA	NA	NA	5	Singly and doubly protonated ions observed based on isotope distribution
15	19.97	0.30	<i>m/z</i> 905.6828 and <i>m/z</i> 453.3455	NA	Unknown	NA	NA	NA	5	Singly and doubly protonated ions observed based on isotope distribution
16 <sup>k</sup>	57.00	0.66	255.2558	35.2	Heptadecylamine	4200-95-7	C <sub>17</sub> H <sub>37</sub> N	NH <sub>2</sub>	3	No CAD response detected for quantitation estimate.
	Total Area (%)	96.97						·		•

<sup>a</sup> Area (%) = (Peak Area / Total Peak Area) × 100.

<sup>b</sup> Reported observed mass-to-charge (*m*/*z*) ratio when peak identification was not definitive (Confidence Levels 4 and 5). <sup>c</sup> Assignment based on published guidelines.<sup>4</sup>

<sup>d</sup> Level 1: Confirmed structure – Matched retention time, accurate mass, and fragmentation pattern to reference standard. <sup>e</sup> Level 2: Proposed structure – Matched accurate mass and fragmentation pattern with library data or published literature; indicates single possible structure.

<sup>f</sup>Level 3: Tentative structure – Matched accurate mass and some fragments yet insufficient evidence to assign single structure (i.e., positional isomers).

<sup>g</sup> Level 4: Unequivocal molecular formula - Unambiguous match to molecular formula without fragmentation information.

<sup>h</sup> Level 5: Exact mass of interest – Accurate mass without sufficient evidence to assign molecular formula or structure.

<sup>i</sup> Low abundance ( $\leq 0.30\%$ ) peak eluting near (D)-Catechin identified as Epicatechin.

<sup>j</sup> Peak 10A/10B are unresolved chromatographic peaks with unique mass spectral profiles; Area % is reported as a total peak area. <sup>k</sup> Peak 16 (Retention Time ~ 57.00 min) is not displayed in partial chromatogram presented in Figure 2.

### **Estimation of Concentration Using CAD Response**

Quantitation of *Ephedra sinica* extract components identified by accurate mass measurements and library searching (i.e., assigned Confidence Level 3; no components were assigned Confidence Level 2) was attempted using CAD responses as an estimate based on (D)-catechin response and concentration (i.e., single-point calibration). The mean CAD peak area ratio of the component to (D)-catechin was multiplied by the (D)-catechin concentration determined by standard addition quantitation (0.351% (w/w); see Table 8). CAD estimated results are summarized in Table 4.

Compound	CAS	Determined Conc. Replicate 1 (% w/w)	Determined Conc. Replicate 2 (% w/w)	Determined Conc. Replicate 3 (% w/w)	Mean ± SD (RSD, %)
Armepavine	524-20-9	0.321	0.270	0.332	0.308 ± 0.033 (10.7%)
Afzelechin/Epi- afzelechinª	2545-00-8 and/or 36801-69-1	0.190	0.193	0.249	0.211 ± 0.033 (15.6%)
Apigenin-5- rhamnoside and Vitexin rhamnoside and Kaempferol-3-O- rhamnoside	Unassigned and 32426-34-9 and Unassigned	0.360	0.355	0.383	0.366 ± 0.015 (4.1%)
Herbacetin/Quercetin	527-95-7 and/or 117-39-5	0.222	0.222	0.243	0.229 ± 0.012 (5.2%)

Table 4. CAD Estimated Results Summary for Ephedra sinica Extract

<sup>a</sup> Coeluting unknown identified by (+) ESI-HRMS may contribute to CAD response, artificially elevating the final result (see Additional Comments, Table 3).

UPLC Parameters:					
Column:	Avantor ACE Excel C18-PFP, 1.7 µm, 100x 2.1 mm (Advanced Chromatography Technologies, UK)				
Autosampler Temperature:	6°C				
Column Temperature:	30°C				
Injection Volume:	1 μL				
Flow Rate:	0.3 mL/min				
Mobile Phase A:	1% (v/v) Acetic Acid in Water				
Mobile Phase B:	1% (v/v) Acetic Acid in Acetonitrile				
Gradient Elution Profiles:	<u>Forward</u> : 0% B held for 0.5 min; ramp to 5% B over 4 min; ramp to 15% B over 9 min; decreased to 0% B in 0.5 min and held for 1 min. <u>Inverse</u> : 100% B held for 0.5 min; decrease to 95% B over 4 min; decrease to 85% B over 9 min; ramped to 100% B in 0.5 min and held for 1 min.				
Run Time:	15 minutes				
Retention Time (min):	Norephedrine: 7.00 Quinoline: 7.30 Norpseudoephedrine: 7.59 (L)-Ephedrine: 9.20 Pseudoephedrine: 9.85 N-Methylephedrine: 10.95				
UV (PDA) Parameters:					
UV (PDA) Parameters: 2D Channel Wavelength:	254 nm				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range:	254 nm 190 to 400 nm, 1.2 nm resolution				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters:	254 nm 190 to 400 nm, 1.2 nm resolution				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode <i>m</i> /z 100 to <i>m</i> /z 1200				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode <i>m</i> /z 100 to <i>m</i> /z 1200 0.3 s				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass:	254 nm         190 to 400 nm, 1.2 nm resolution         10 points/sec         1000         35°C         20 pA/1.0         57.5 psi         ESI+; MSe sensitivity mode         m/z 100 to m/z 1200         0.3 s         Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s)				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode <i>m</i> /z 100 to <i>m</i> /z 1200 0.3 s Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s) 2.5 kV				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode <i>m</i> /z 100 to <i>m</i> /z 1200 0.3 s Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s) 2.5 kV 40 V				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage: Collision Energy:	254 nm 190 to 400 nm, 1.2 nm resolution 10 points/sec 1000 35°C 20 pA/1.0 57.5 psi ESI+; MSe sensitivity mode <i>m</i> /z 100 to <i>m</i> /z 1200 0.3 s Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s) 2.5 kV 40 V Low CE: 6 eV High CE: 20-30 eV				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage: Collision Energy: Source Temperature:	254 nm         190 to 400 nm, 1.2 nm resolution         10 points/sec         1000         35°C         20 pA/1.0         57.5 psi         ESI+; MSe sensitivity mode         m/z 100 to m/z 1200         0.3 s         Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s)         2.5 kV         40 V         Low CE: 6 eV       High CE: 20-30 eV         110°C				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage: Cone Voltage: Collision Energy: Source Temperature: Desolvation Temperature:	254 nm         190 to 400 nm, 1.2 nm resolution         10 points/sec         1000         35°C         20 pA/1.0         57.5 psi         ESI+; MSe sensitivity mode         m/z 100 to m/z 1200         0.3 s         Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s)         2.5 kV         40 V         Low CE: 6 eV       High CE: 20-30 eV         110°C         250°C				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage: Cone Voltage: Cone Voltage: Collision Energy: Source Temperature: Desolvation Temperature: Cone Gas Flow:	254 nm         190 to 400 nm, 1.2 nm resolution         10 points/sec         1000         35°C         20 pA/1.0         57.5 psi         ESI+; MSe sensitivity mode         m/z 100 to m/z 1200         0.3 s         Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s)         2.5 kV         40 V         Low CE: 6 eV       High CE: 20-30 eV         110°C         250°C         50 L/h				
UV (PDA) Parameters: 2D Channel Wavelength: 3D Channel Wavelength Range: CAD Parameters: Sampling Rate: Scale Factor: Evaporation Temperature: Scale/Power Function Value: Nitrogen HRMS Parameters: Ionization/Acquisition Mode: Acquisition Range: Scan Time: Lock Mass: Capillary Voltage: Cone Voltage: Cone Voltage: Collision Energy: Source Temperature: Desolvation Temperature: Cone Gas Flow: Desolvation Gas Flow:	254 nm         190 to 400 nm, 1.2 nm resolution         10 points/sec         1000         35°C         20 pA/1.0         57.5 psi         ESI+; MSe sensitivity mode         m/z 100 to m/z 1200         0.3 s         Leucine encephalin (m/z 278.1135, 556.2766; Scan for 0.300 s, interval: 30 s)         2.5 kV         40 V         Low CE: 6 eV       High CE: 20-30 eV         110°C         250°C         50 L/h         600 L/h				

 Table 5. Standard Addition Quantitation - Acquisition Parameters

# **Standard Addition Quantitation**

Standard addition calibration solutions were prepared as presented in Figure 1. Individual stock standard solutions of ephedrine hydrochloride (as (L)-ephedrine), pseudoephedrine hydrochloride (as pseudoephedrine), DL-norephedrine hydrochloride (as norephedrine), norpseudoephedrine, N-methylephedrine, (D)-catechin, quinoline, and 1-phenylpropane-1,2-dione were prepared at ~ 1 mg/mL, diluted in 95% ethanol. Mixed component spiking solutions were prepared at three concentrations in 50% (v/v) aqueous ethanol (1.0, 2.5, and 5.0  $\mu$ g/mL for each component) for quantitation by standard addition methodology.

The linear regression equation relating peak areas of the standard addition calibration solutions to their additional analyte concentration was determined for data without weighting. A 4-point calibration curve with triplicate standards at the lower (unspiked) and upper limits of the concentration range was used for quantitation. The concentration of each analyte in the test article solution was calculated by extrapolation of the linear regression equation to the x-axis, application of appropriate dilution factor(s), and unit conversion to express results as a weight-to-weight percentage.

Representative extracted ion chromatograms, standard addition calibration curves, and quantitation summary tables are presented in Figures 3 - 12 and Tables 6 - 12.



**Figure 3.** Extracted ion chromatograms (m/z 134.0964) for diluent blank (black), unspiked extract (A: Set A and B: Set C; red), and spiked extract (~ 250 ng/mL added; blue).



Figure 4. Extracted ion chromatograms (m/z 148.1126) for diluent blank (black), unspiked extract (Set B; red), and spiked extract (~ 100 ng/mL added; blue).



**Figure 5.** Extracted ion chromatograms (m/z 180.1383) for diluent blank (black), unspiked extract (Set A; red), and spiked extract (~ 100 ng/mL added; blue).



**Figure 6.** Extracted ion chromatograms (m/z 291.0863) for diluent blank (black), unspiked extract (Set A; red), and spiked extract (~ 100 ng/mL added; blue).

#### **Standard Addition Quantitation – Calibration**



Figure 7. Representative standard addition calibration curve - Norephedrine

Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Intercept <sup>a</sup>	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	45.104	177947	0.6379	3945.3	0.198
BSC-EtOH-2	10.486	28.108	169511	0.5935	6030.8	0.320
BSC-EtOH-3	10.276	32.949	155967	0.5393	4733.6	0.256
					Mean ± SD	0.258 ± 0.061
					(% RSD)	(23.6%)

**Table 6.** Norephedrine Quantitative Results Summary for *Ephedra sinica* Extract

<sup>a</sup> X-Intercept = Absolute value of (Y-Intercept/Slope).

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/450  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.



Figure 8. Representative standard addition calibration curve – Norpseudoephedrine

<b>Fable 7.</b> Norps	eudoephedrine	Quantitative Result	s Summary for	r <i>Ephedra sinica</i> I	Extract
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Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Intercept <sup>a</sup>	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	81.521	13194	0.9887	161.85	0.366
BSC-EtOH-2	10.486	77.741	12398	0.9795	159.47	0.380
BSC-EtOH-3	10.276	73.963	12049	0.9773	162.91	0.396
					Mean ± SD	0.381 ± 0.015
					(% RSD)	(3.9%)

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/10  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.



Figure 9. Representative standard addition calibration curve – (L)-Ephedrine

Table 8. (L)	-Ephedrine	Quantitative	Results	Summary	for	Ephedra	sinica	Extract
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Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Interceptª	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	129.00	13494	0.9988	104.60	2.36
BSC-EtOH-2	10.486	111.14	12223	0.9939	109.97	2.62
BSC-EtOH-3	10.276	99.776	11849	0.9956	118.76	2.89
					Mean ± SD	2.62 ± 0.27
					(% RSD)	(10.1%)

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/50  $\mu$ L) × (500  $\mu$ L/10  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.



Figure 10. Representative standard addition calibration curve – Pseudoephedrine

Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Intercept <sup>a</sup>	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	138.80	6269.2	0.9949	45.166	1.02
BSC-EtOH-2	10.486	121.91	7804.4	0.9895	64.015	1.53
BSC-EtOH-3	10.276	113.62	5776.4	0.9927	50.837	1.24
					Mean ± SD	1.26 ± 0.26
					(% RSD)	(20.2%)

**Table 9.** Pseudophedrine Quantitative Results Summary for *Ephedra sinica* Extract

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/50  $\mu$ L) × (500  $\mu$ L/10  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.



Figure 11. Representative standard addition calibration curve – N-Methylephedrine

<b>Table 10.</b> N-Methylephedrine	Quantitative Results Summar	y for <i>Ephedra sinica</i> Extract
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Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Intercept <sup>a</sup>	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	2093.2	148051	0.9989	70.729	0.160
BSC-EtOH-2	10.486	2080.9	130116	0.9922	62.528	0.149
BSC-EtOH-3	10.276	1851.6	129970	0.9990	70.192	0.171
					Mean ± SD	0.160 ± 0.011
					(% RSD)	(6.9%)

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/10  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.



Figure 12. Representative standard addition calibration curve – (D)-Catechin

Extract Replicate	Subsample (mg)	Slope	Y-Intercept	Correlation Coefficient (R <sup>2</sup> )	X- Interceptª	Determined Concentration <sup>b</sup> (% w/w)
BSC-EtOH-1	11.058	184.53	29700	0.9968	160.95	0.364
BSC-EtOH-2	10.486	182.49	23847	0.9828	130.67	0.312
BSC-EtOH-3	10.276	157.51	24467	0.9960	155.34	0.378
					Mean ± SD	0.351 ± 0.035
					(% RSD)	(10.0%)

<sup>b</sup> Determined Conc. (% w/w) = X-Intercept × (500  $\mu$ L/10  $\mu$ L) × (5 mL/2.5 mL) × (2.5 mL/Subsample in mg) × (1000 mg/1 g) × (1 g/10<sup>9</sup> ng) × 100.

Reference Compound	CAS	Determined Conc. Replicate 1 (% w/w)	Determined Conc. Replicate 2 (% w/w)	Determined Conc. Replicate 3 (% w/w)	Mean ± SD (RSD, %)
Norephedrine	14838-15-4	0.198	0.320	0.256	0.258 ± 0.061 (23.6%)
Quinoline	91-22-5	ND	ND	ND	ND
Norpseudoephedrine	492-39-7	0.366	0.380	0.396	0.381 ± 0.015 (3.9%)
(L)-Ephedrine	299-42-3	2.36	2.62	2.89	2.62 ± 0.27 (10.1%)
Pseudoephedrine	90-82-4	1.02	1.53	1.24	1.26 ± 0.26 (20.2%)
N-Methylephedrine <sup>a</sup>	552-79-4	0.160	0.149	0.171	0.160 ± 0.011 (6.9%)
(D)-Catechin	154-23-4	0.364	0.312	0.378	0.351 ± 0.035 (10.0%)

Table 12. Standard Addition Quantitative Results Summary for *Ephedra sinica* Extract

ND = No response detected above baseline noise in undiluted extract solutions. <sup>a</sup> Possible coelution with pseudomethylephedrine.