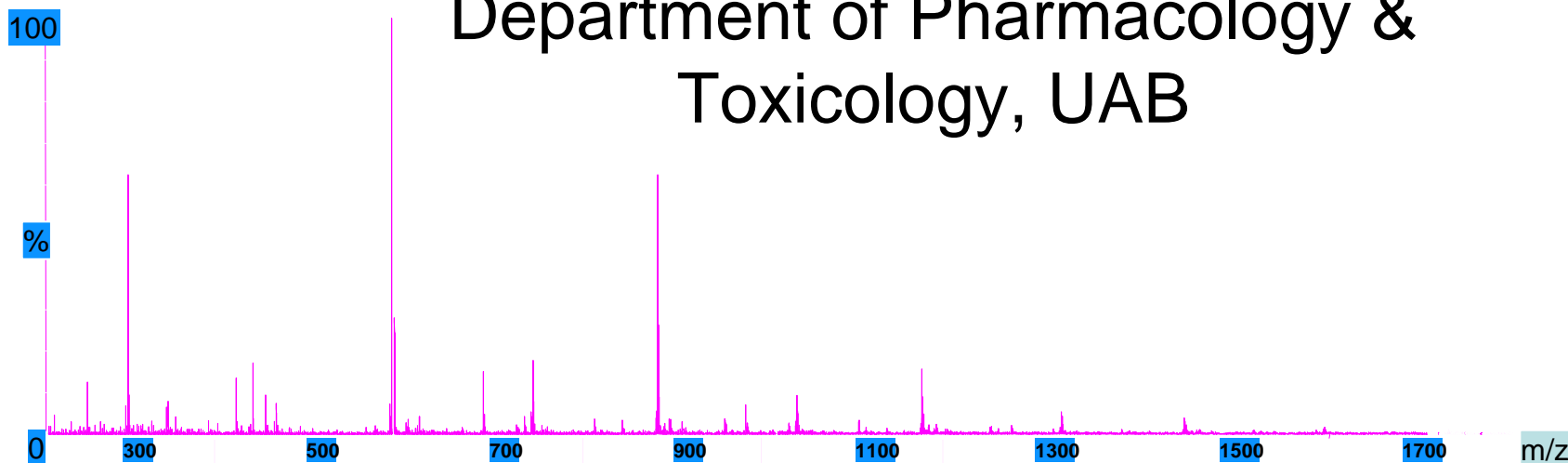


# Qualitative and quantitative analysis/method validation in metabolomics

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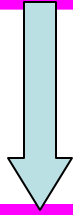


# **Class Overview**

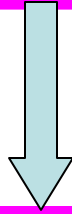
- **Introduction to bioanalysis**
- **Quantitative analysis of puerarin, and isoflavones in biological samples by LC-MS/MS**

# ***Bioanalysis Flow Chart***

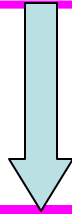
**Sample preparation**



**Chromatographic separation**

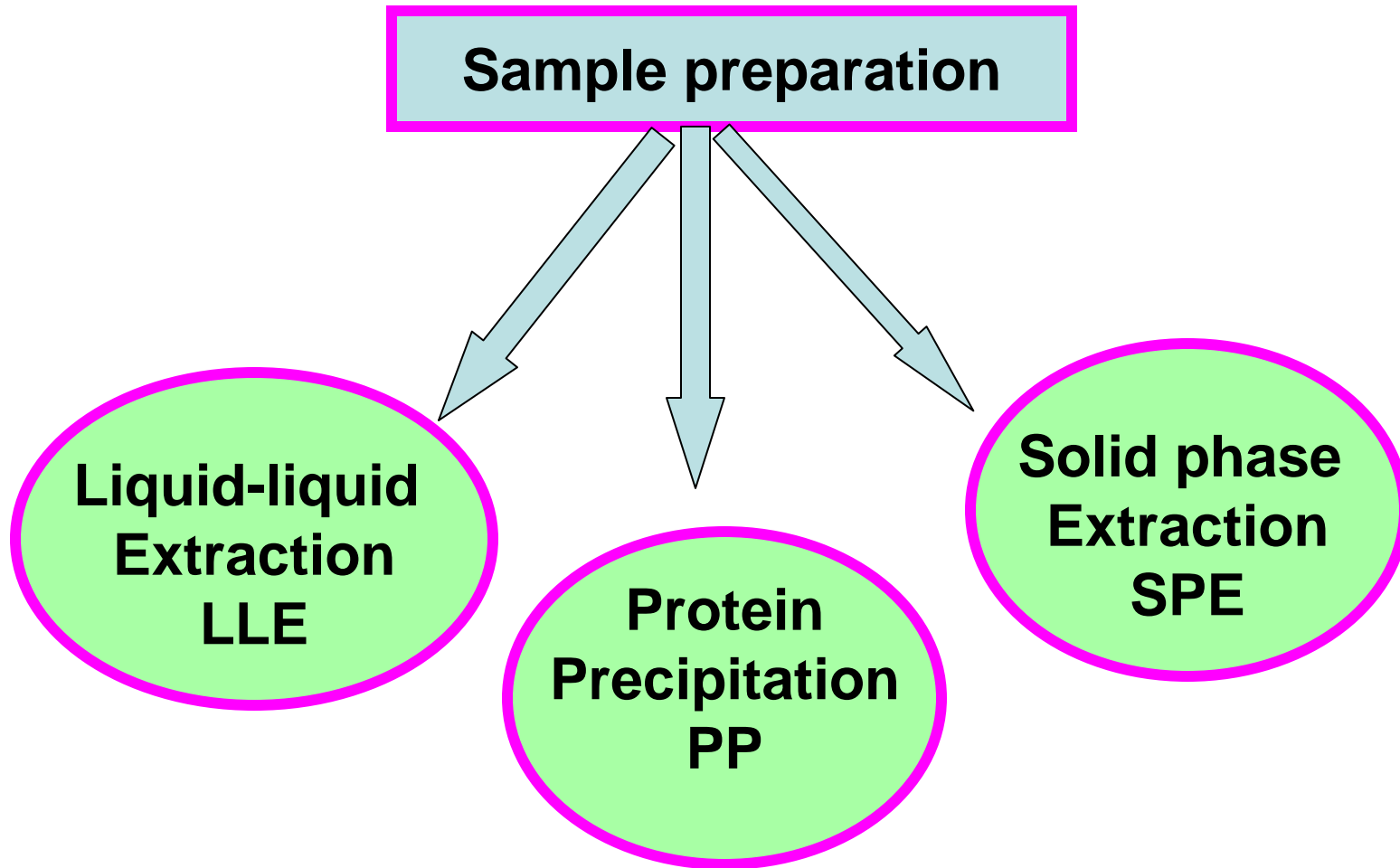


**MS ionization**



**MS analysis**

***Sample preparation is a crucial step in removing the interfering compounds from biological matrix***



***The method of choice will be determined by the sample matrix and the concentration of compounds in samples***

# **Choice of Good Internal Standards**

- **A stable isotopically labeled IS is preferable.**
- **Is not found in the original sample**
- **In the absence of stable isotopically labeled internal std, the structure of the internal standard needs to be similar to the analyte and co-elute with the analyte.**
- **Should not react chemically with the analyte.**

# Factors affecting ionization of analytes

- Polarity and ionization potential of analytes (ESI vs APCI)
- Mobile phase composition (appropriate use of buffer such as acids and base), methanol vs acetonitrile, water composition in elution etc.
- Matrix components/salts
- Ion source of hardware

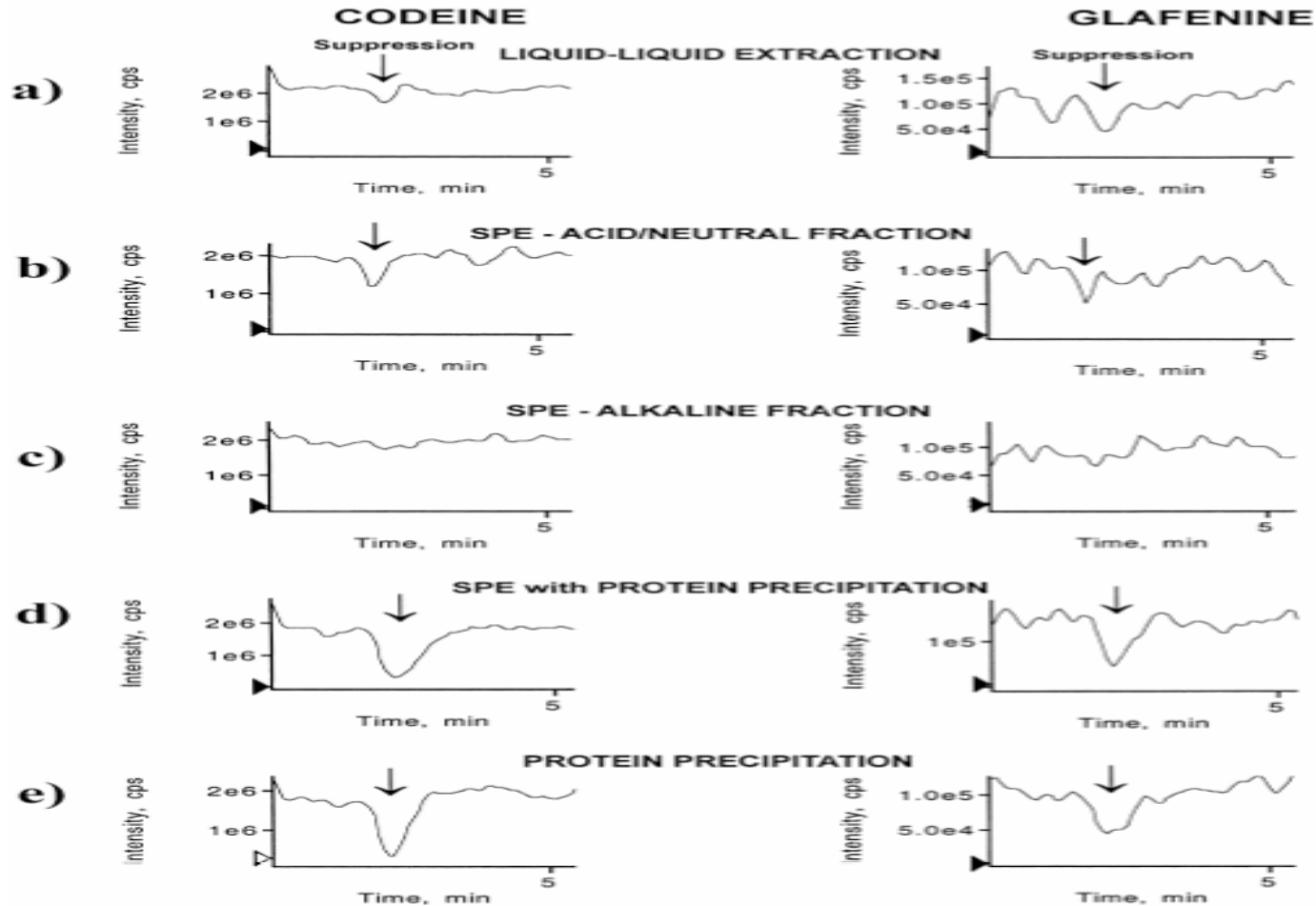
# Problems encountered in LC-MS analysis

## Matrix effect

### Ion suppression?

- the presence of endogenous substances from matrix, i.e. organic or inorganic molecules present in the sample and that are retrieved in the final extract
- exogenous substances, i.e. molecules not present in the sample but coming from various external sources during the sample preparation

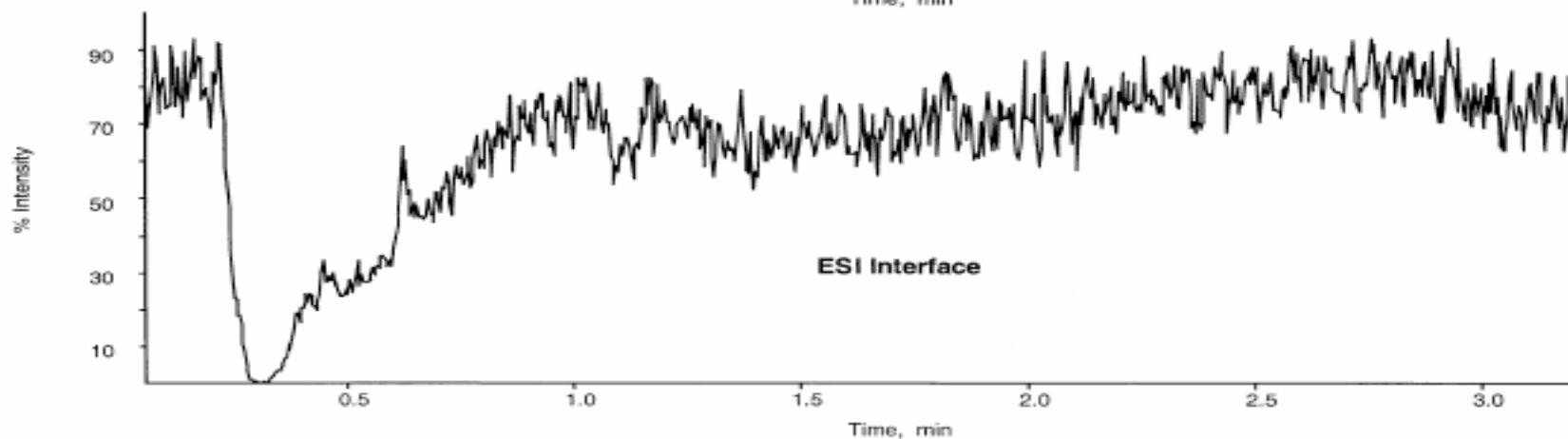
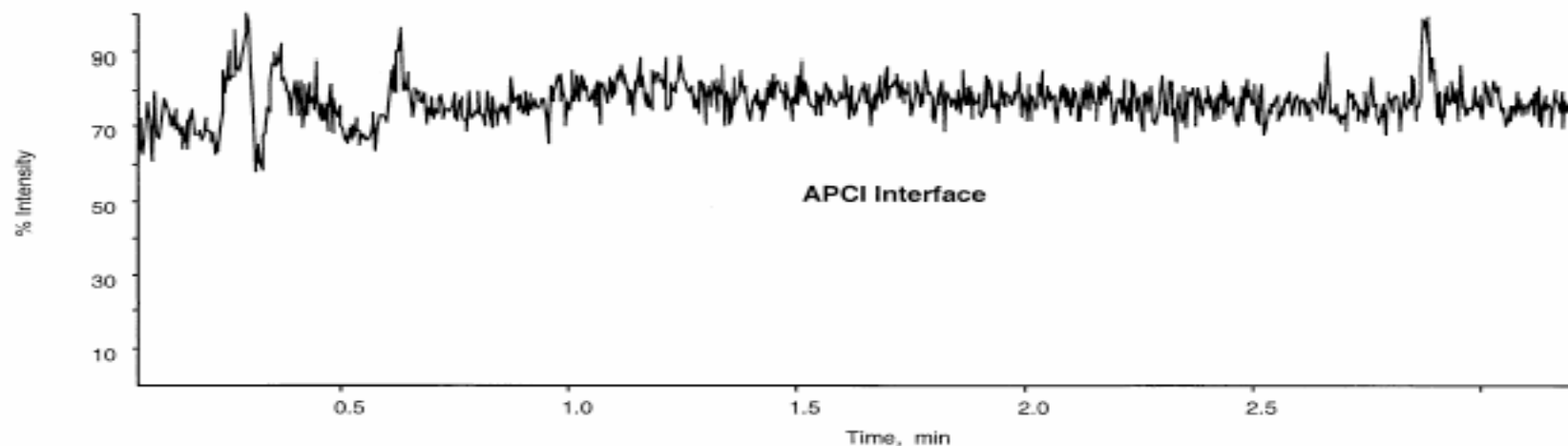
# Severe ion suppression effect for codein and glafenin was observed with PPT and SPE-PPT



Muller et al. *J. Chrom B* (2002)



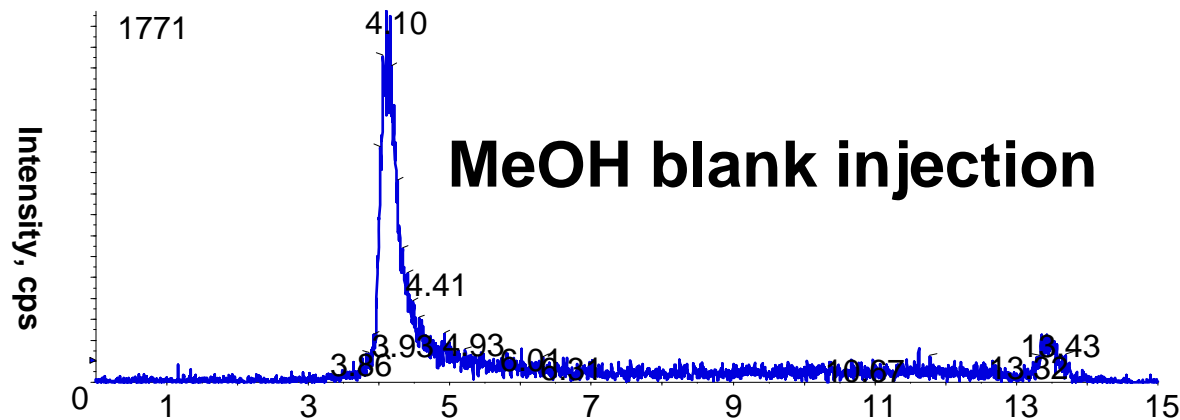
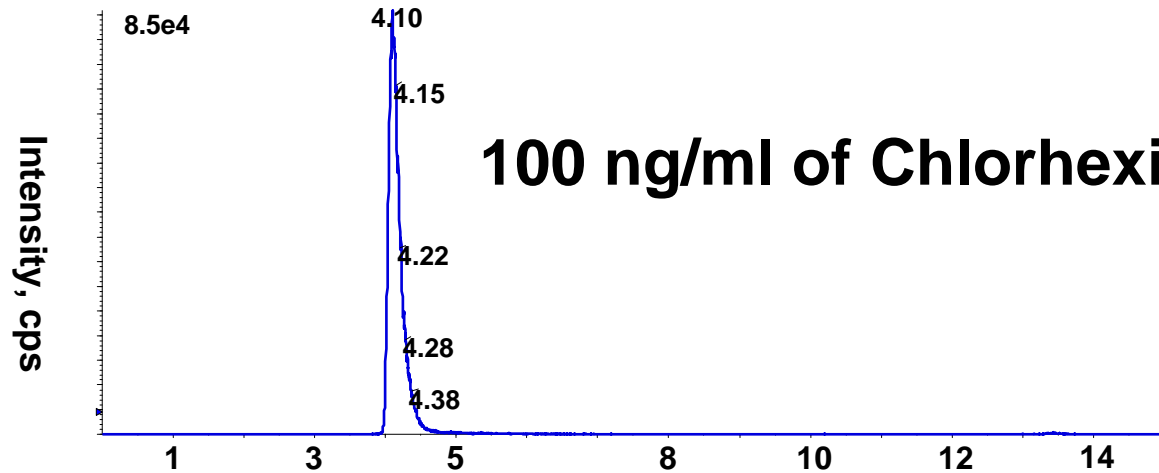
# APCI is less prone to than ESI to the effects of ion suppression



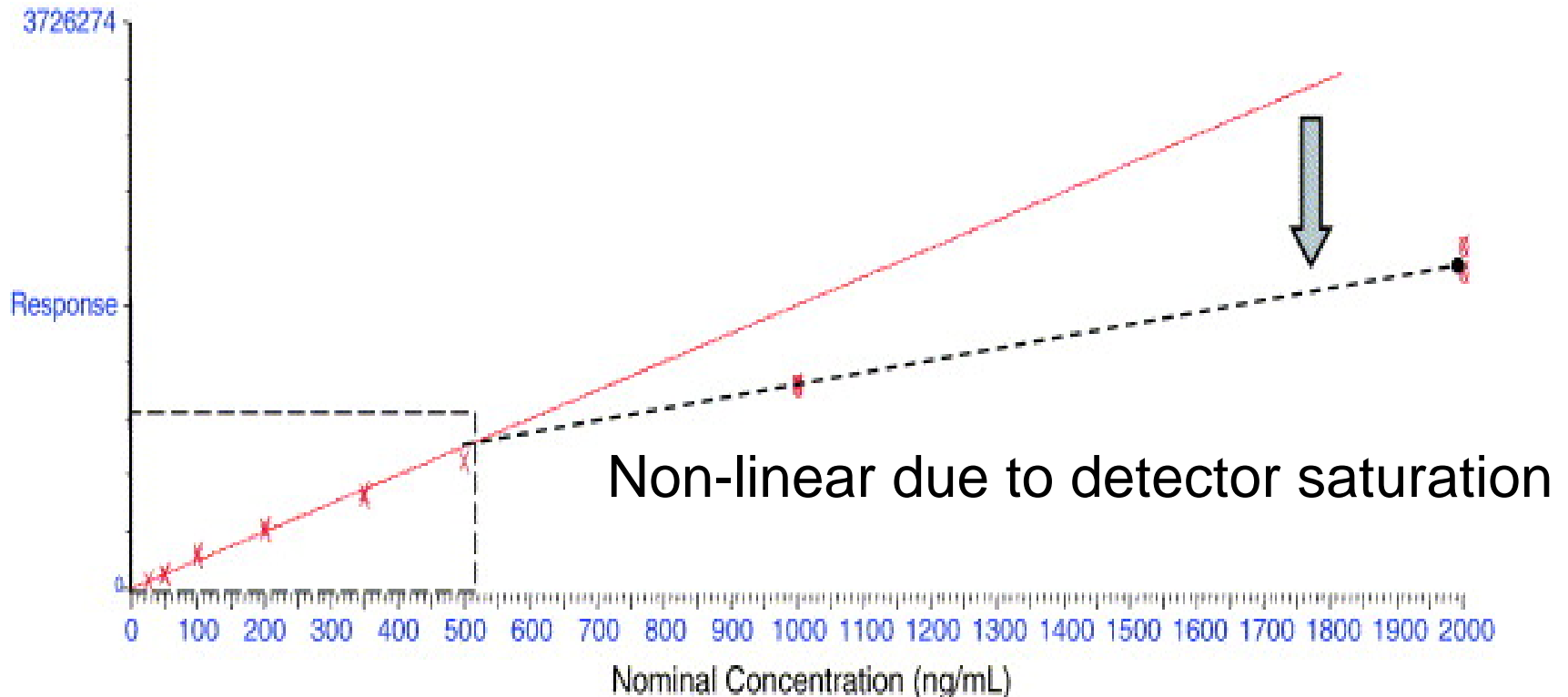
*King et al. J. Am Soc Mass Spectrom 2000*

# Carry over a big problem?

Previously injected sample when appears upon subsequent analyses due to physico-chemical property of the sample, analysis system or both.



# Standard curve non-linearity is possible due to Detector saturation, dimer/multimer formation, and or ESI droplet saturation at higher concentration



Source: Bakhtiar & Majumdar.

Journal of Pharmacological and Toxicological methods, 2007

# **Why quantification of drug/drug metabolites in plasma/tissues PK studies is so important?**

- **An accurate and fast analytical method for measuring the concentrations of a compound in plasma or tissue is the first step in order to yield the PK of a compound**
- **Established assay for human sample analyses (plasma, serum or urine matrix) needs to be more rugged, robust and be able to withstand the test of time during this the longest phase of clinical development. The requirements and adherence to specificity, selectivity and stability will become very important**

# **Analytical method validation**

- **Should demonstrate specificity, linearity, recovery, accuracy, precision**
- **Lower limit of quantification**
- **Stability (freeze/thaw)**
- **Robustness**
- **Matrix effects**

# Method validation..

- **Specificity is established by the lack of interference peaks at the retention time for the internal standard and the analyte.**
- **Accuracy is determined by comparing the calculated concentration using calibration curves to known concentration. The LLQ is defined as the smallest amount of the analyte that could be measured in a sample with sufficient precision (%CV) and accuracy (within 20% for both parameters) and is chosen as the lowest concentration on the calibration curve.**

# LC/MS/MS Method for Puerarin

**Column:** Waters X-Terra C18 with guard,  
2.1 x 100 mm, 3.5 micron

**Mobile Phase A:** 10% MeCN + 10 mM NH<sub>4</sub>OAc

**Mobile Phase B:** 70% MeCN + 10mM NH<sub>4</sub>OAc

**Gradient:** 0 minutes = 100% A

6 minutes = 100% B

7 minutes = 100% A

10 minutes = Stop

**Injection Volume:** 20 ul

**Flow Rate:** 0.2 ml/min split flow

**Mass Spectrometer:** Negative Electrospray

**Mass Transitions:** 415/267 (Puerarin)

415/295 (Puerarin)

269/149 (apigenin, IS)

Table 1.  
Summary of calibration curves (n =5)

Concentration (ng/ml)	Mean $\pm$ S.D.	CV (%)	Accuracy (%)
2.0	2.21 $\pm$ 0.16	7.00	110.7
5.0	5.22 $\pm$ 0.28	5.30	104.48
50	45.32 $\pm$ 2.53	5.60	90.64
500	473.60 $\pm$ 26.57	5.60	94.72
1000	1021.20 $\pm$ 71.53	7.00	102.12
5000	5340 $\pm$ 420.18	7.90	106.80

Mean r = 0.996

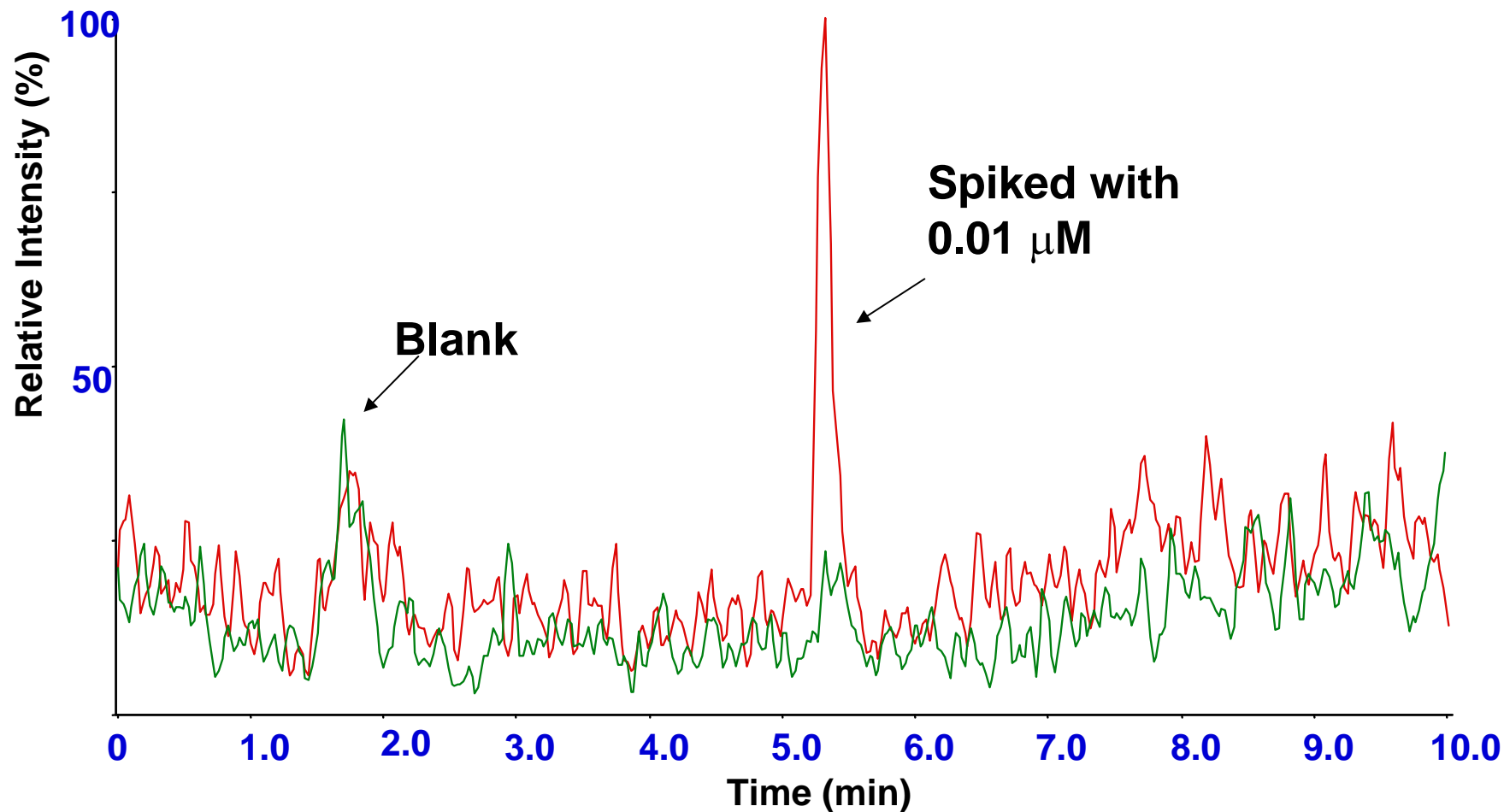


Table 2.

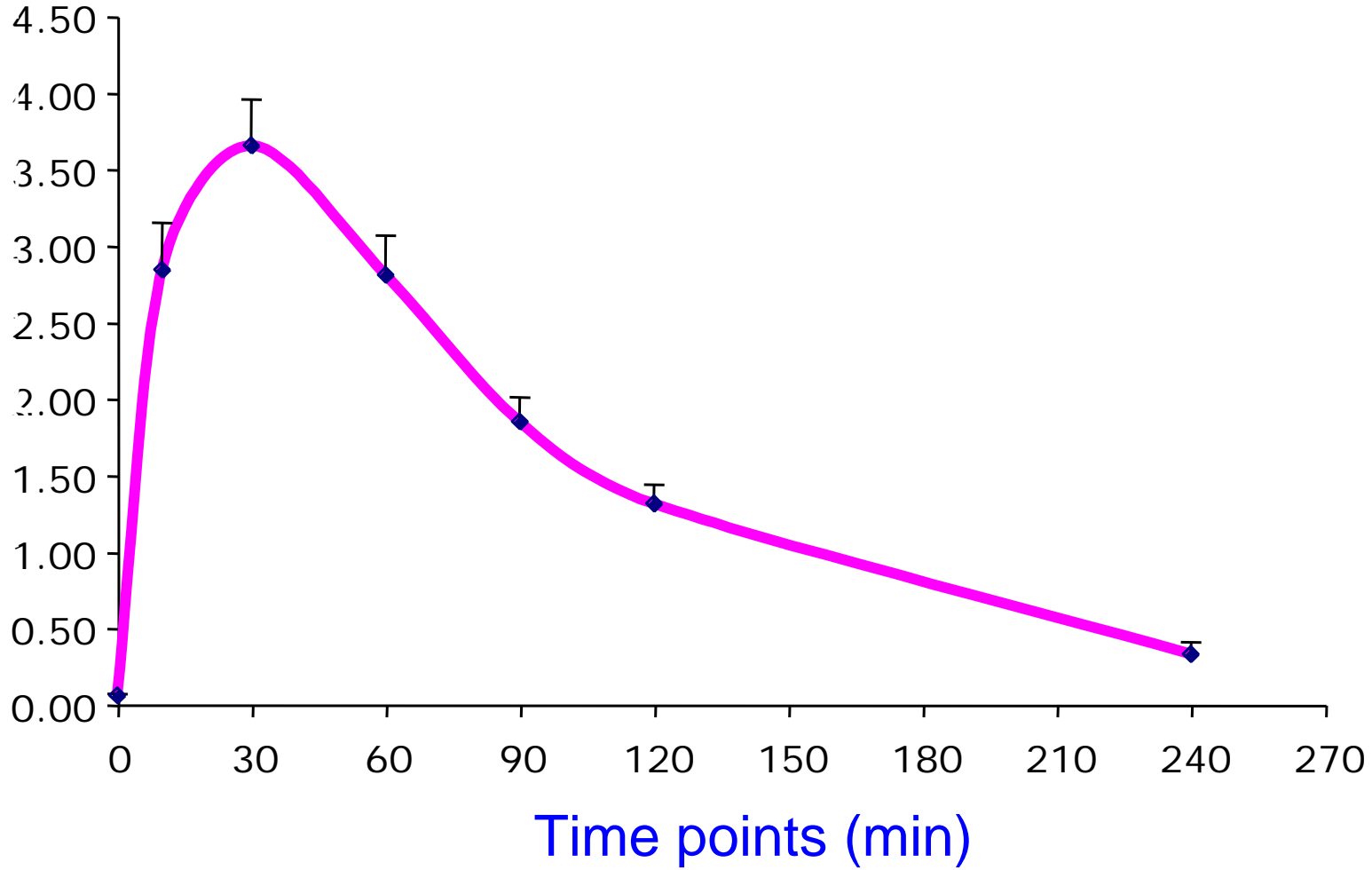
Assay validation characteristics of the method for the determination of puerarin in rat serum (n =5)

Concentration (ng/ml)	Mean $\pm$ S.D.	CV (%)	Accuracy (%)
2.0	2.21 $\pm$ 0.16	7.00	110.7
4.0	3.96 $\pm$ 0.30	7.90	99.20
8.32	7.32 $\pm$ 1.00	14.40	113.30
20	19.20 $\pm$ 1.20	6.30	96.00
200	203.20 $\pm$ 19.41	9.60	101.60
832	821.18 $\pm$ 55.86	6.80	101.31
2000	2240 $\pm$ 96.70	4.30	112.00

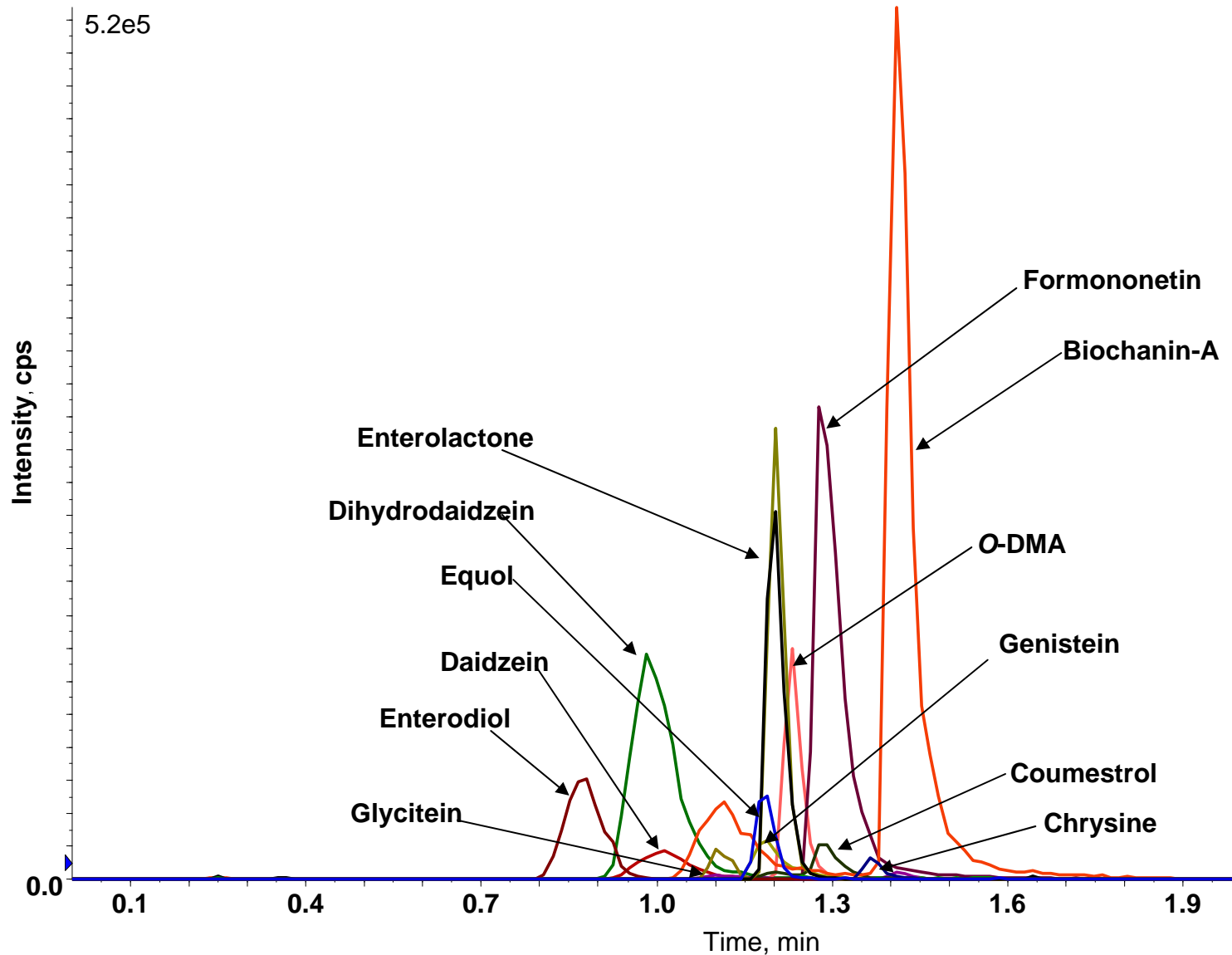
# Ion chromatograms of a rat serum spiked sample (0.01 $\mu\text{M}$ of puerarin) vs. blank serum



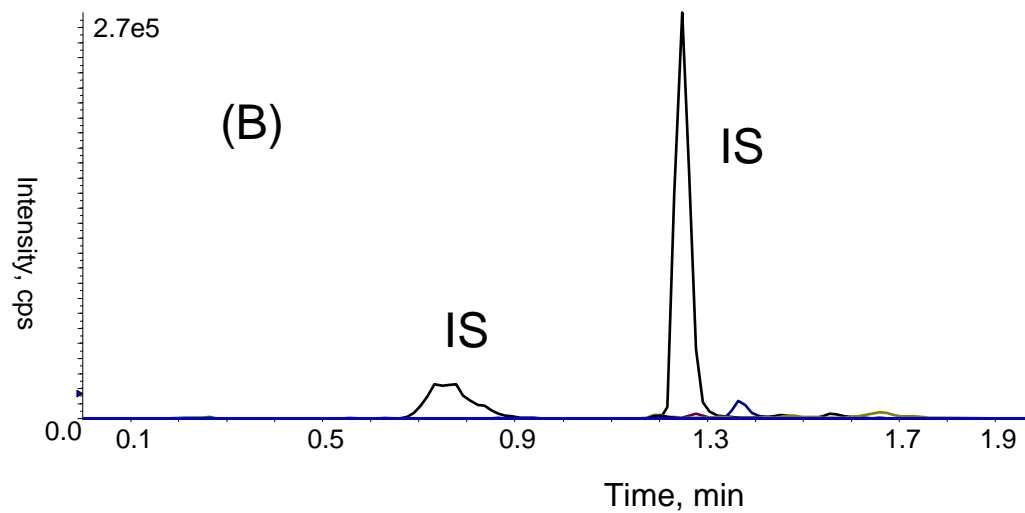
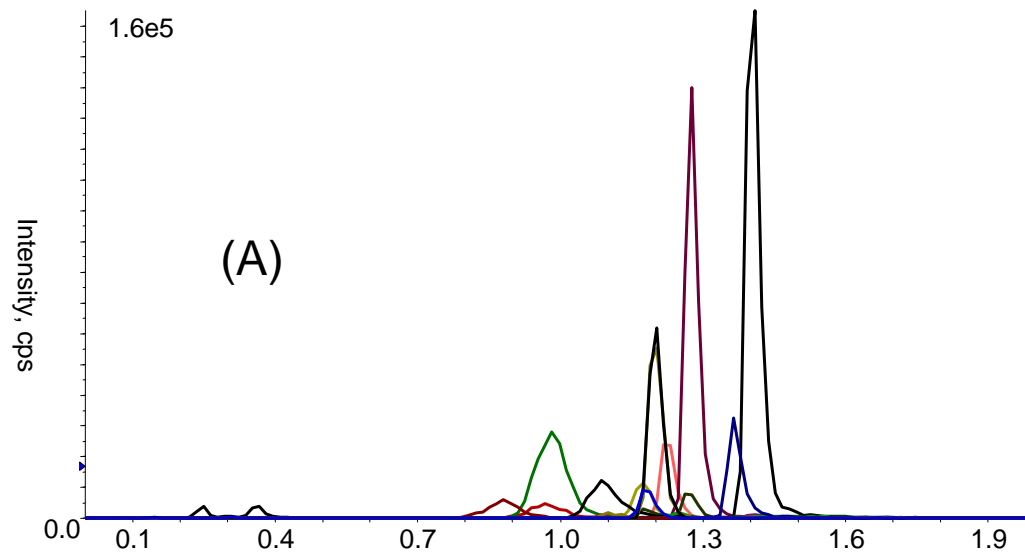
***Average serum concentration of puerarin versus time after Oral administration of 50 mg/kg puerarin***



# MRM chromatogram showing separation of 11 phytoestrogens using 2 min run time



# Specificity of the assay- no peaks from matrix



# Calibration range and lower limit of Quantification (LLOQ) of analytes

Analyte	Calibration range (ng/ml)	LLOQ (ng/ml)
Equol	1 - 5,000	1
Daidzein	2 - 5,000	2
DHD	2 - 5,000	2
O-DMA	1 - 5,000	1
genistein	2 - 5,000	2
Glycitein	5 - 5,000	5
Formononetin	1 - 5,000	1
Coumestrol	1 - 5,000	1
Bichanin-A	1 - 5,000	1
6-OH-ODMA	20 - 5,000	20
Enterodiol	2 - 5,000	2
Enterolactone	1 - 5,000	1

# Precision and accuracy of standards

Analyte	Concentration (ng/mL)	CV (%)	Mean accuracy (%)
Equol	1	9.26	95.82
	2	4.94	99.73
	20	1.86	100.00
	100	4.85	100.05
	200	3.33	106.32
	1000	4.79	100.72
	5000	2.38	98.47
Daidzein	1	6.84	101.56
	2	5.10	99.00
	20	6.83	100.08
	100	1.00	104.50
	200	5.92	101.48
	1000	5.86	99.48
	5000	3.67	96.90
Dihydrodaidzein	1	ND	ND
	2	13.49	97.62
	20	2.06	101.30
	100	5.13	104.05
	200	3.15	107.66
	1000	4.85	102.92
	5000	2.02	99.03
O-DMA	1	6.28	95.53
	2	6.15	96.53
	20	5.00	102.98
	100	2.47	101.55
	200	3.39	103.73
	1000	7.63	103.98
	5000	6.59	96.70
Genistein	1	10.47	104.16
	2	10.73	94.47
	20	6.23	99.18
	100	2.60	104.16
	200	2.72	107.66
	1000	6.72	106.34
	5000	4.62	102.38
Glycitein	1	3.08	114.83
	2	5.92	107.21
	20	7.94	93.91
	100	6.78	100.35
	200	4.10	105.93
	1000	2.90	95.82
	5000	6.21	102.05
Formononetin	1	4.39	100.72
	2	7.40	102.77
	20	8.52	101.18
	100	5.48	97.03

# Precision and accuracy of quality control samples

Analyte	Nominal concentration (ng/mL)	Accuracy (%)			Precision (%CV)			
		Day 1	Day 2	Day 3	Day 1	Day 2	Day 3	Inter-day
Equol	50	100.42	90.13	96.60	2.01	4.33	5.11	3.74
	500	103.30	99.85	114.66	2.31	5.61	1.93	2.97
	2,000	97.60	89.90	103.96	6.11	10.61	10.13	8.34
Daidzein	50	99.98	102.73	94.04	4.35	6.44	8.23	6.62
	500	101.48	98.31	97.73	3.14	5.44	7.42	5.38
	2,000	92.50	87.41	86.03	2.88	3.61	3.96	3.58
Dihydrodaidzein	50	103.00	100.15	101.66	3.94	1.43	4.99	3.63
	500	103.79	95.20	106.00	3.96	6.44	3.35	4.34
	2,000	91.70	90.40	96.33	1.68	5.80	6.60	2.82
O-DMA	50	104.00	93.72	96.51	5.16	4.71	5.80	5.32
	500	105.67	93.78	102.33	3.22	9.42	5.54	5.84
	2,000	101.20	93.57	100.93	5.53	5.37	6.53	3.63
Genistein	50	107.66	106.83	99.08	3.97	3.37	6.65	4.86
	500	97.50	88.90	91.36	5.40	3.61	5.60	4.96
	2,000	95.13	92.28	93.38	2.63	3.97	4.17	3.59