



# **The Science Required For Successful Bioanalytical Methods**

**By: Ridha Nachi**

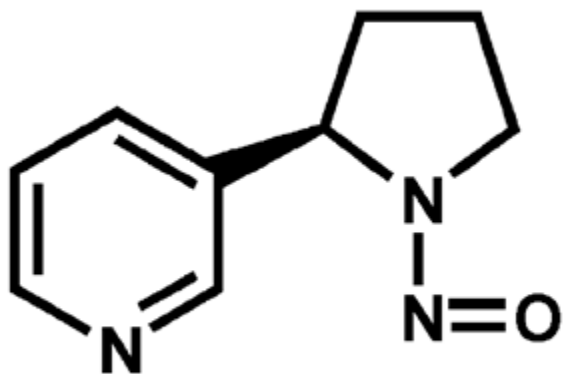
**TSRC, 2012**

# Scope Of Presentation

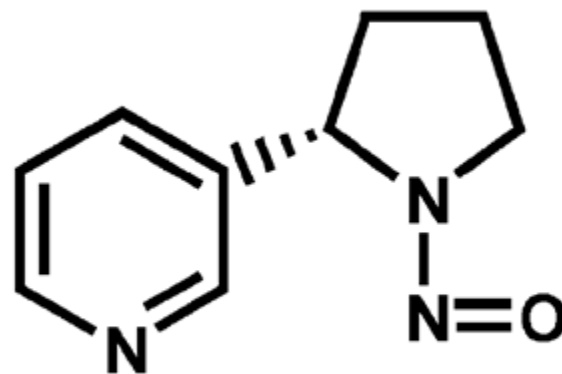
- Selectivity
- Sensitivity
- Matrix effect and accuracy. Are stable-labeled internal standards equivalent?

# *N'*-nitrosornicotine (NNN) Chemical Structures

Target LLOQ 0.75 pg/mL in human urine



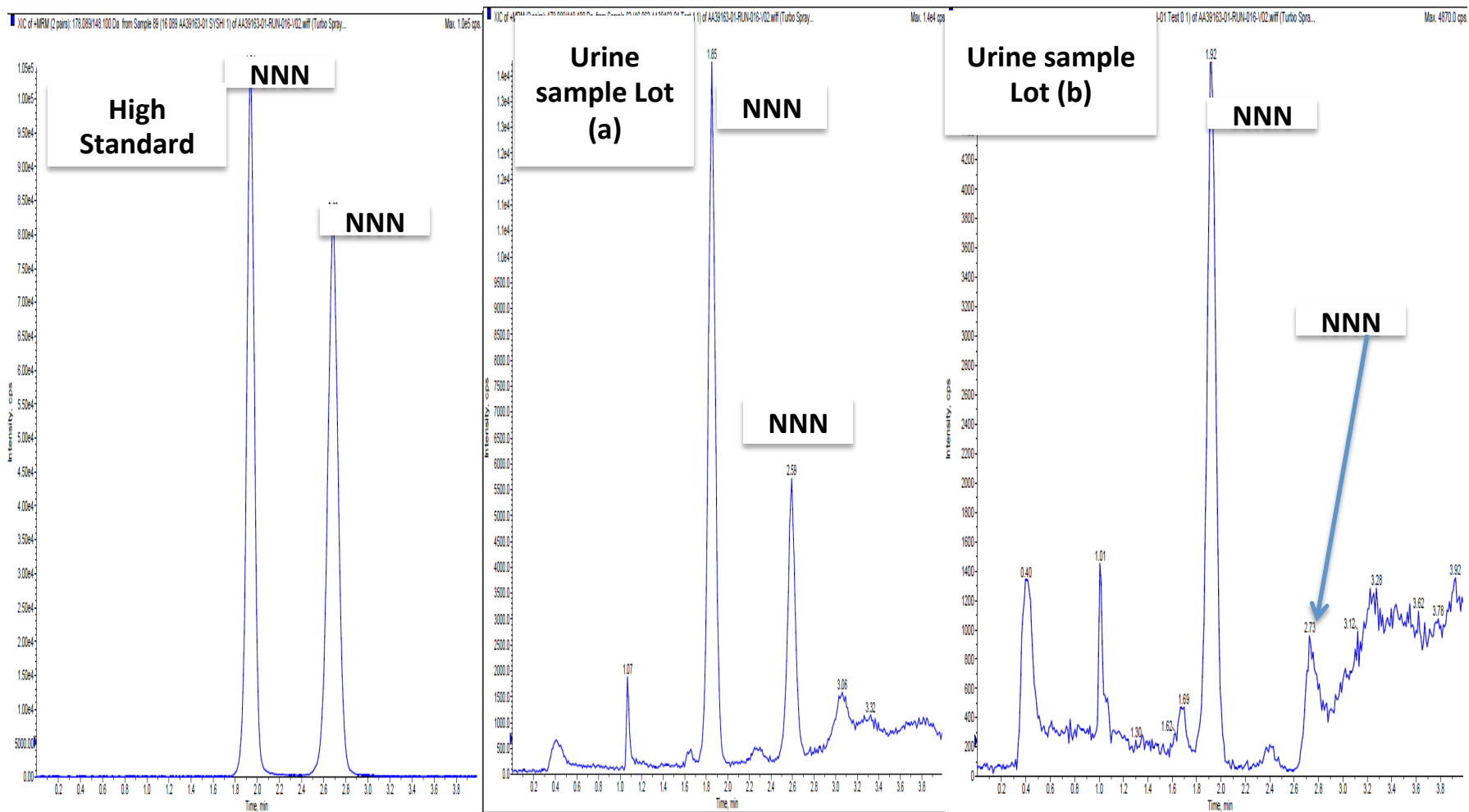
**(R)-NNN**



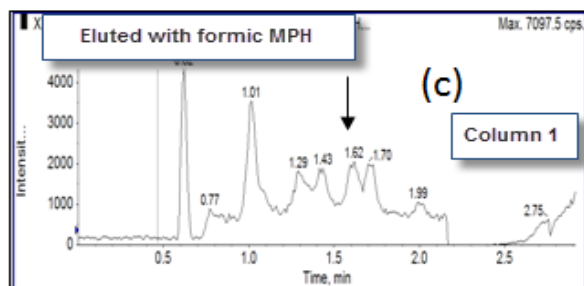
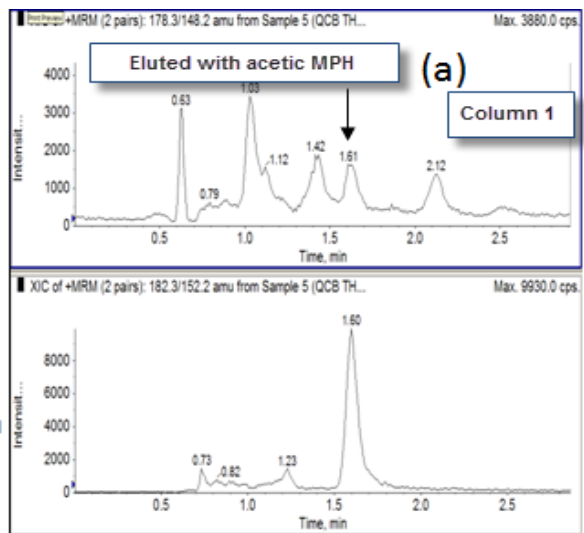
**(S)-NNN**

# NNN Extracted Urine Sample

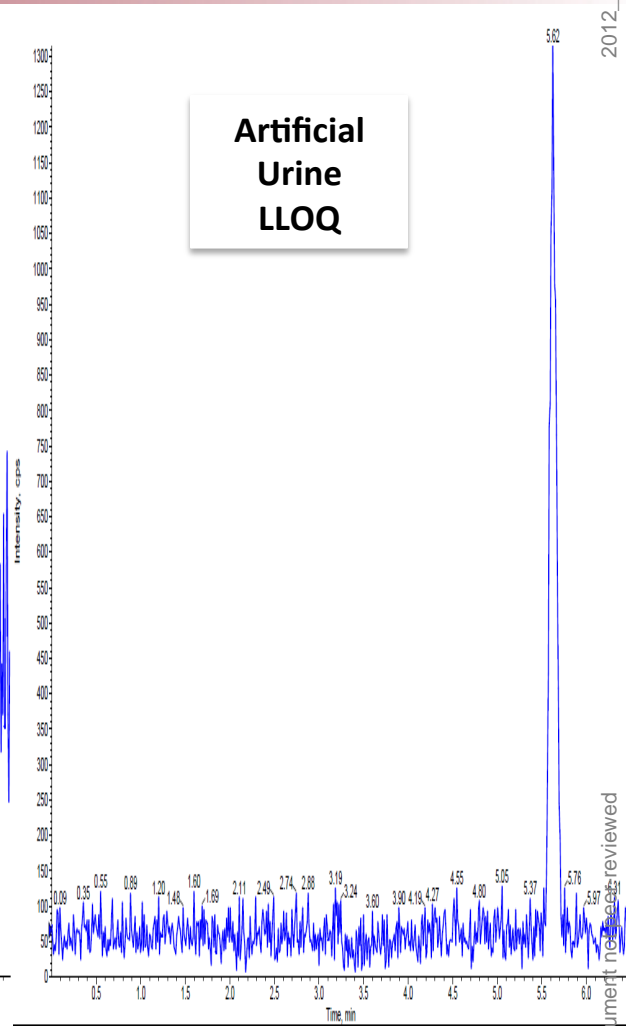
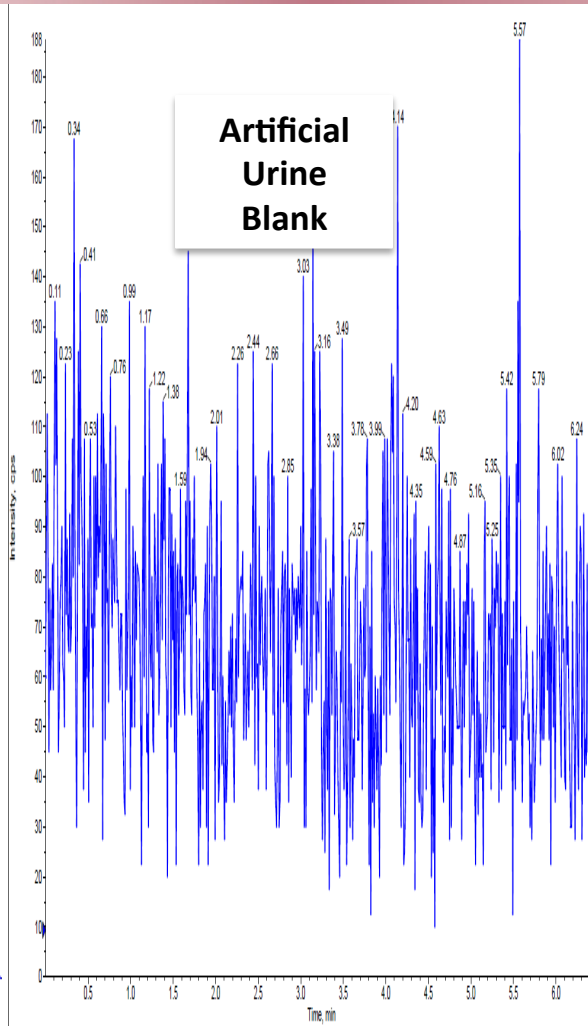
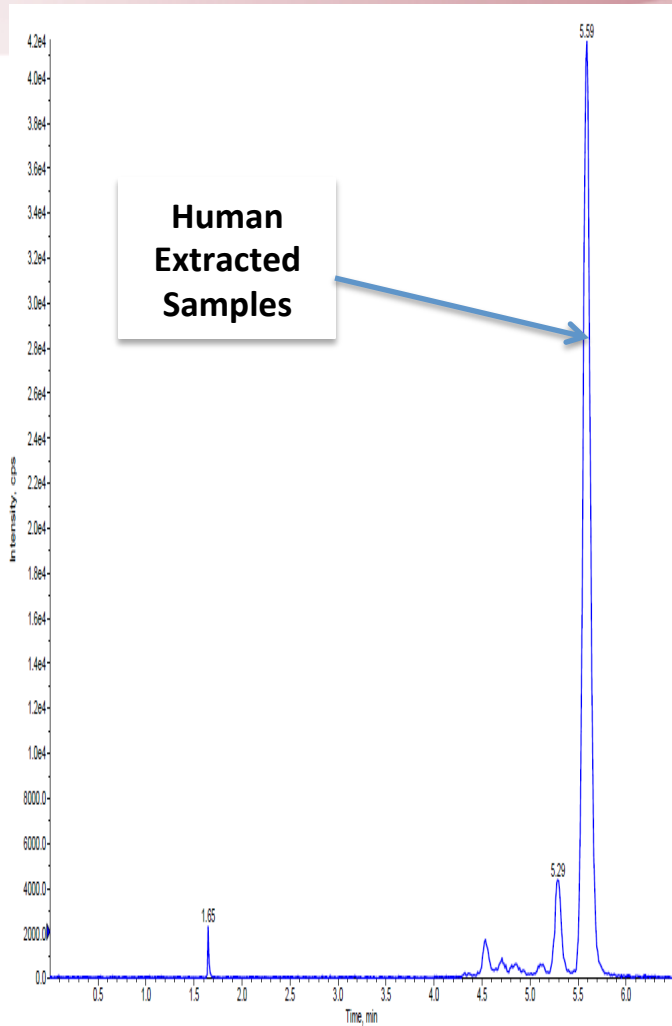
Chromatography on a cation-exchange column (with separation of NNN stereoisomers)



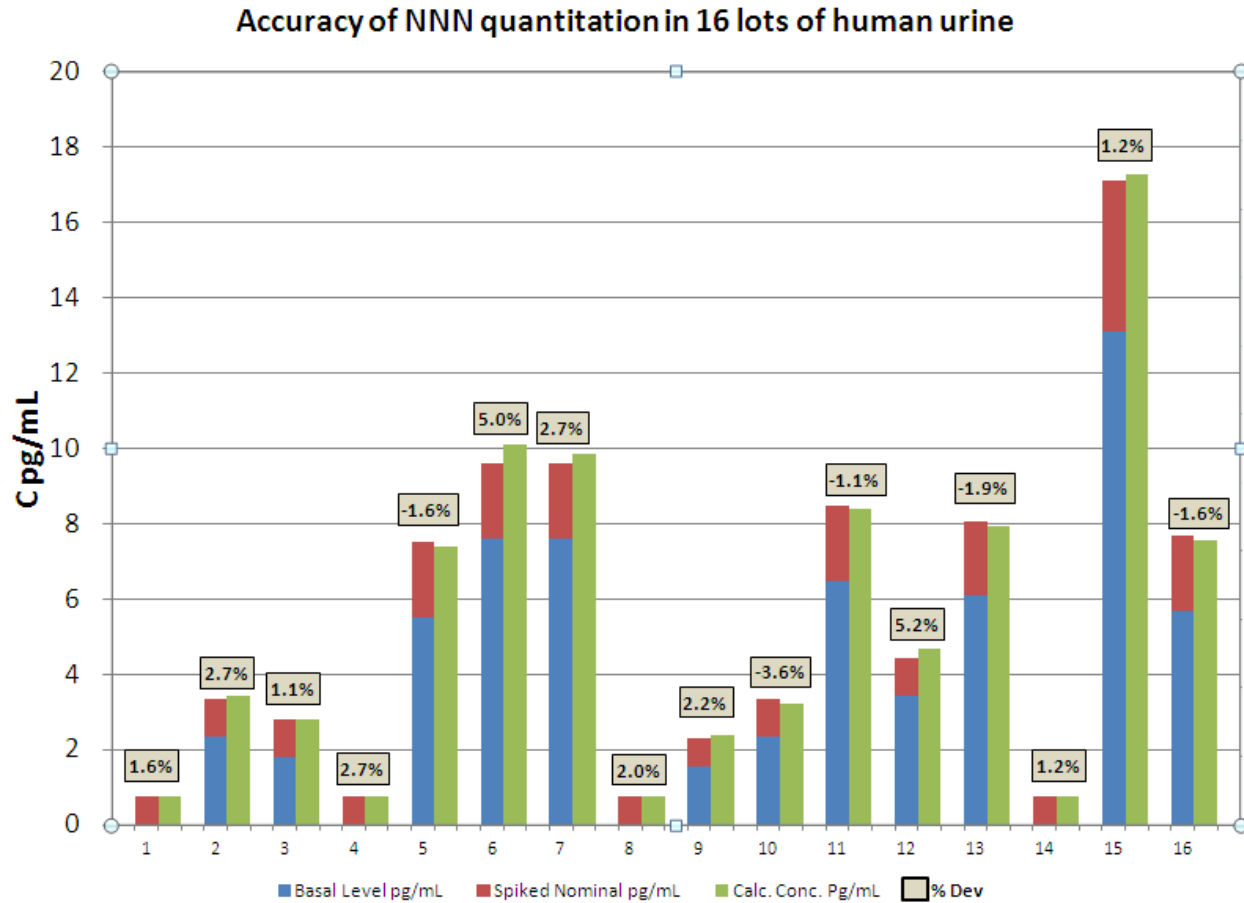
# Two-dimensional chromatography of extracted urine samples on a Polar RP columns



# NNN Urine Extracted LLOQ (0.75pg/mL)

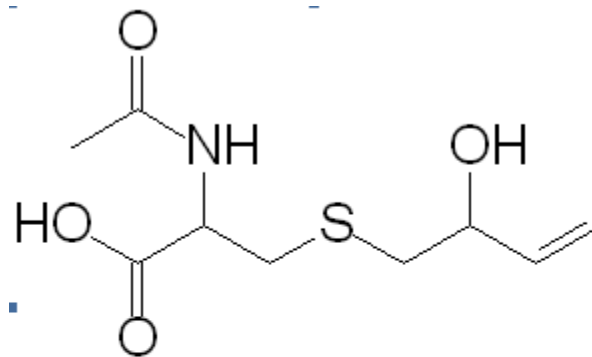


# Accuracy of NNN Quantitation in Human Urine



# Monohydroxy-3-Butenyl Mercapturic Acid (MHBMA) In Human Urine

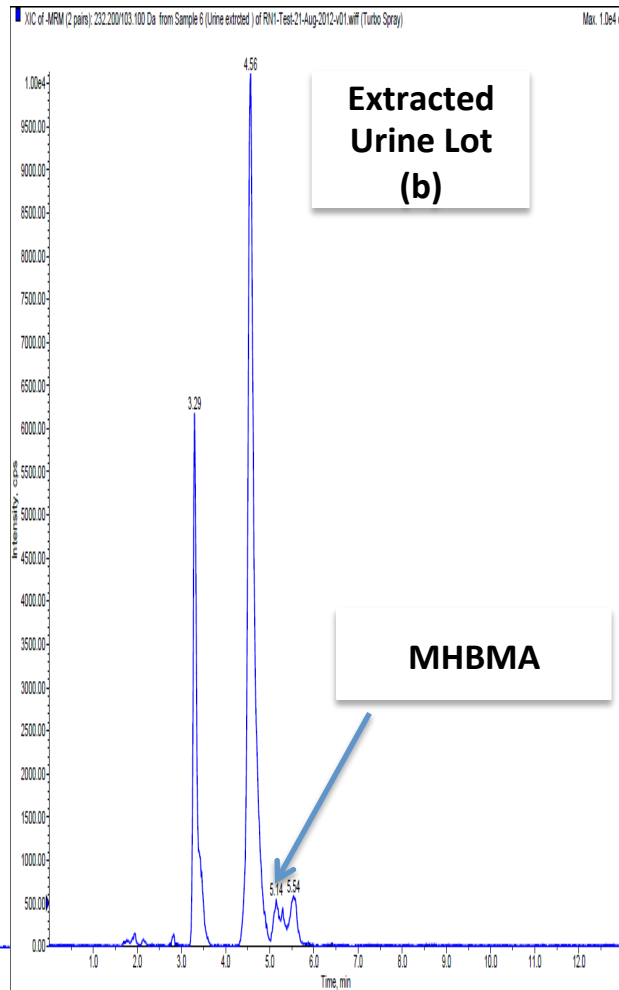
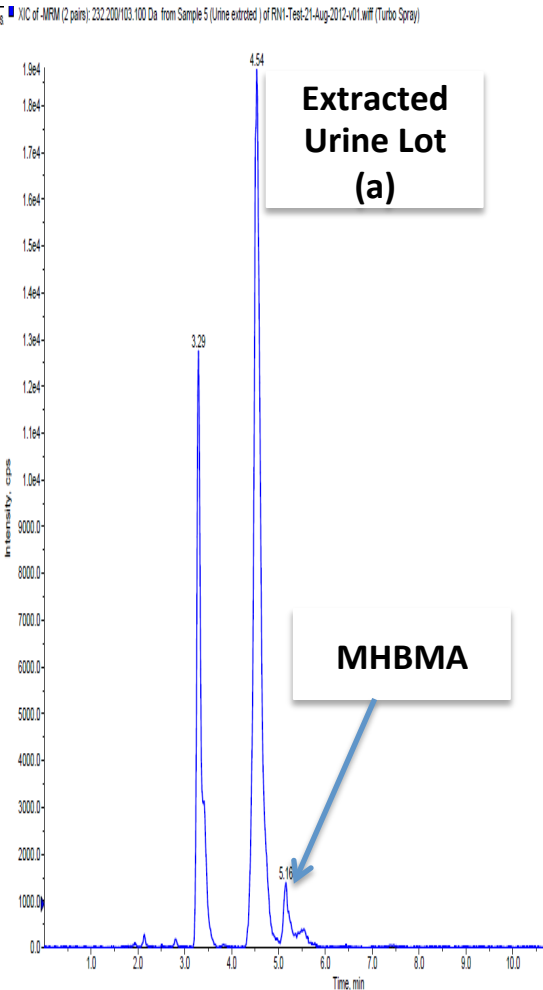
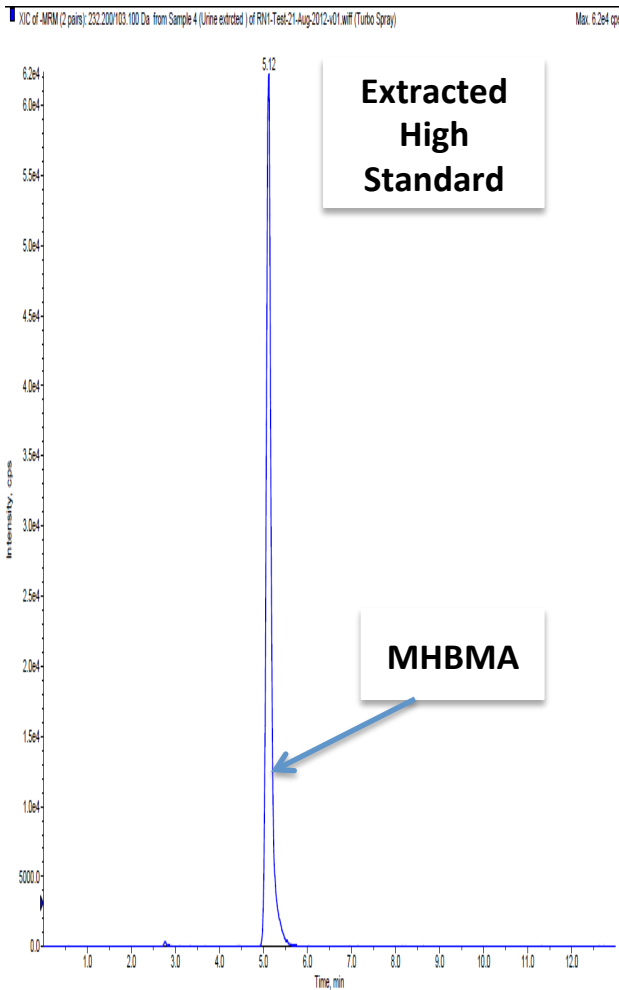
Mixed mode SPE.  
Anion exchange chromatography.  
ESI Negative Ion mode ionization.



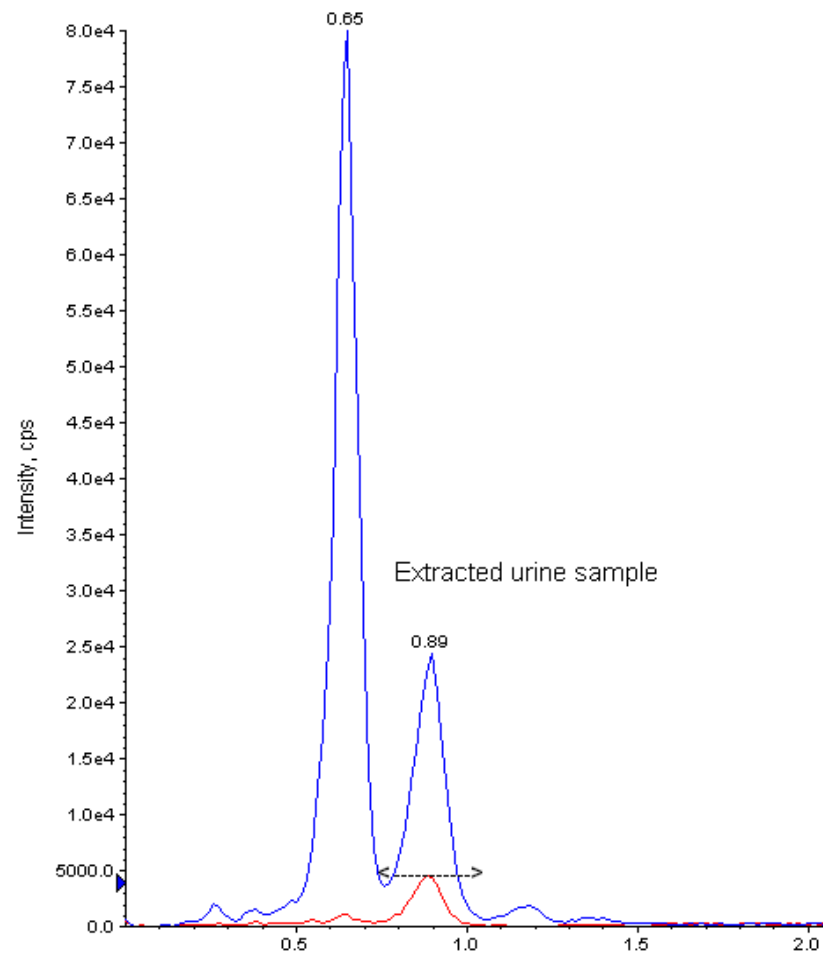
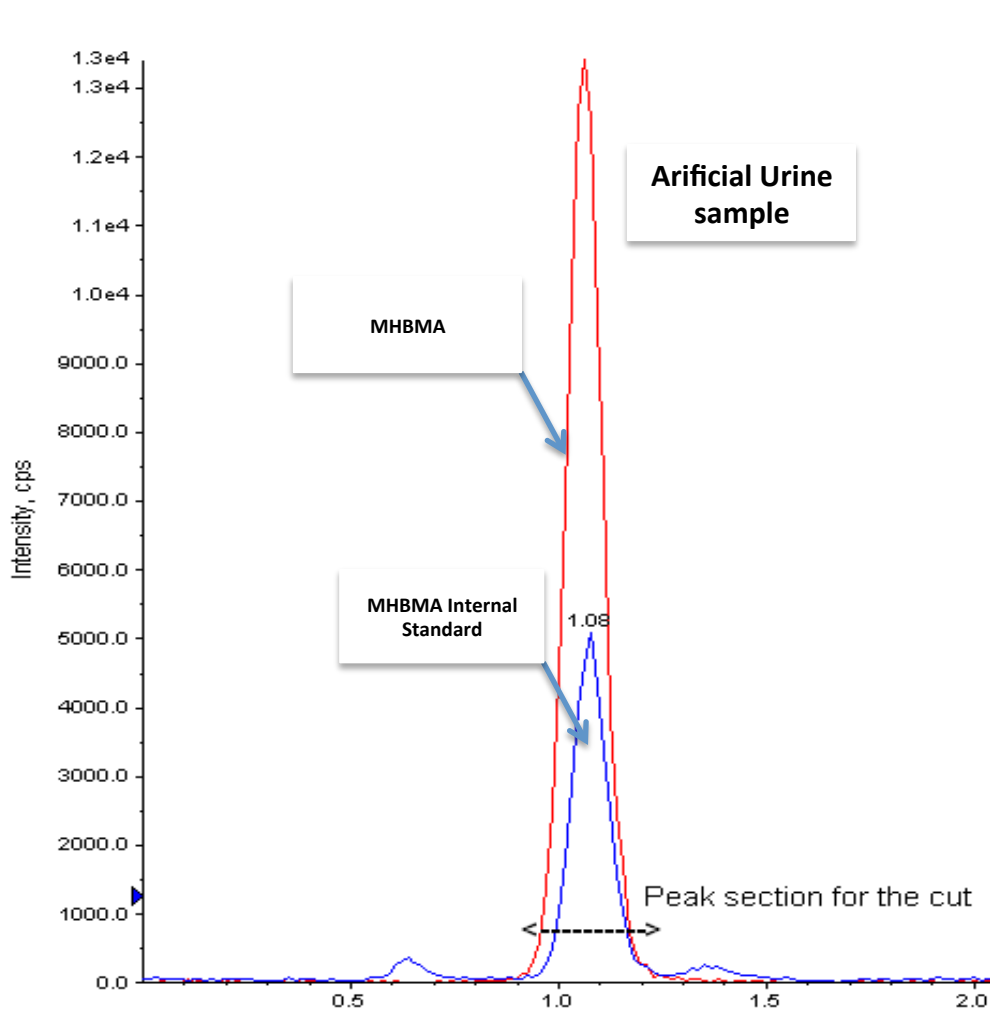
MHBMA  
 $C_9H_{15}NO_4S$   
232.28 Da



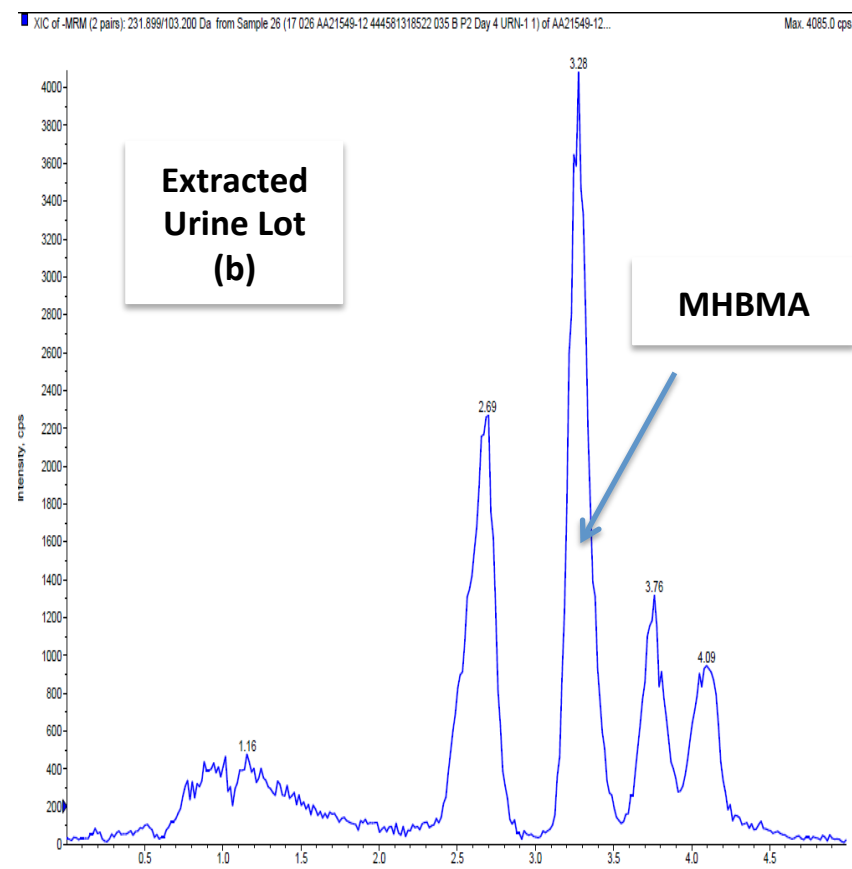
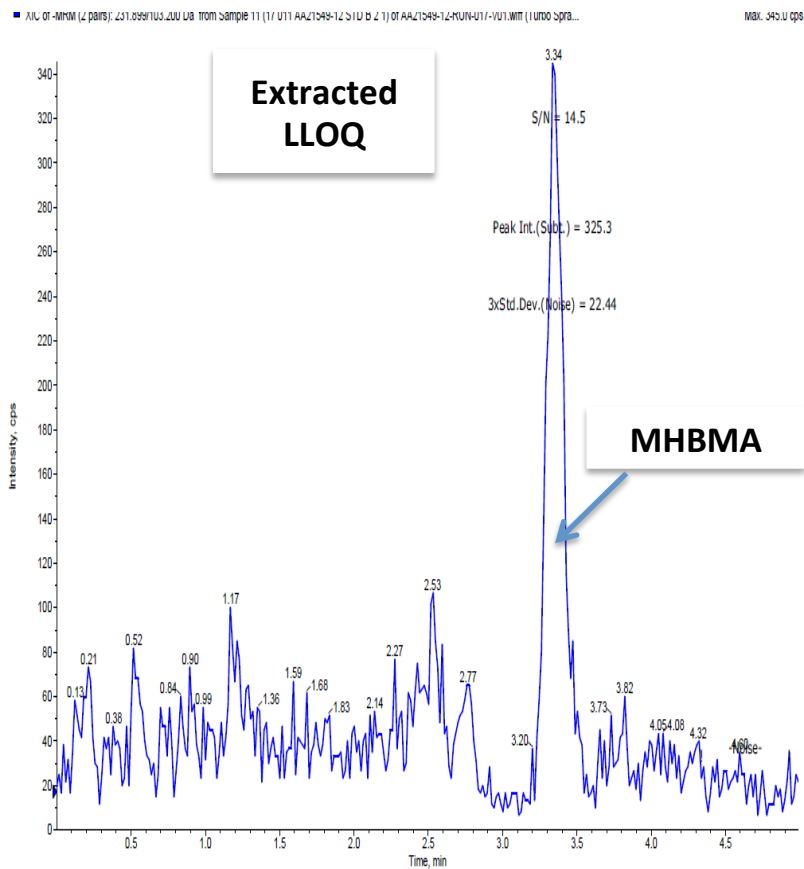
# MHBMA Extracted Samples



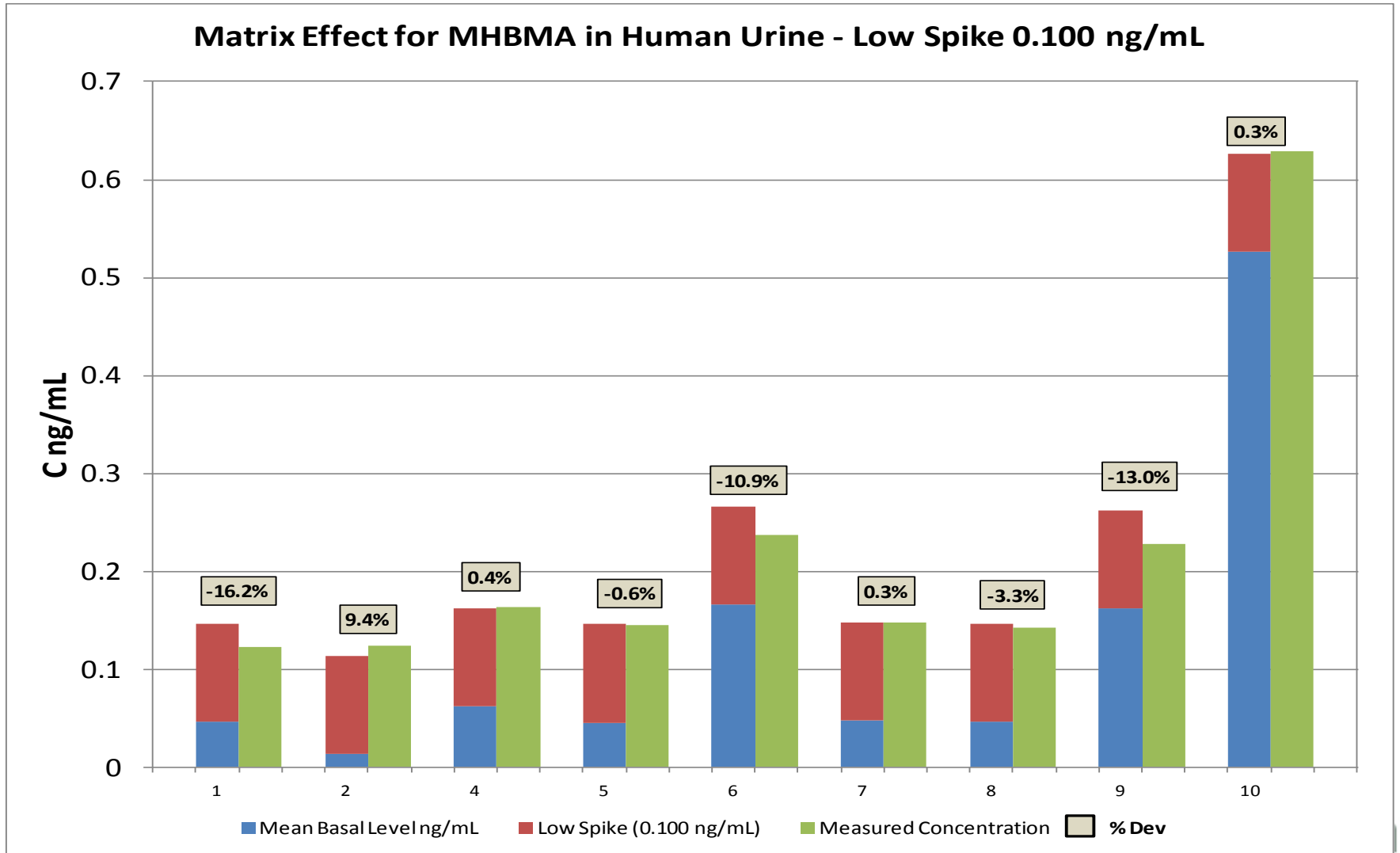
# MHBMA heart cut on an anion exchange guard column



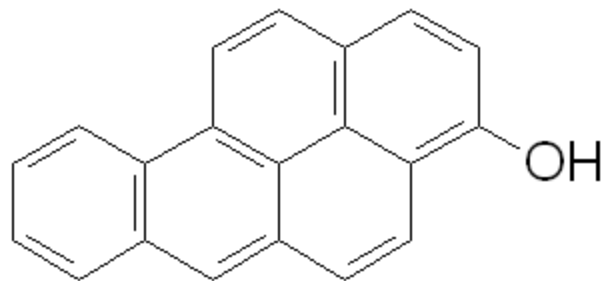
# Examples of MHBMA Final Chromatography



# Accuracy of Matrix Effect for MHBMA in Human Urine

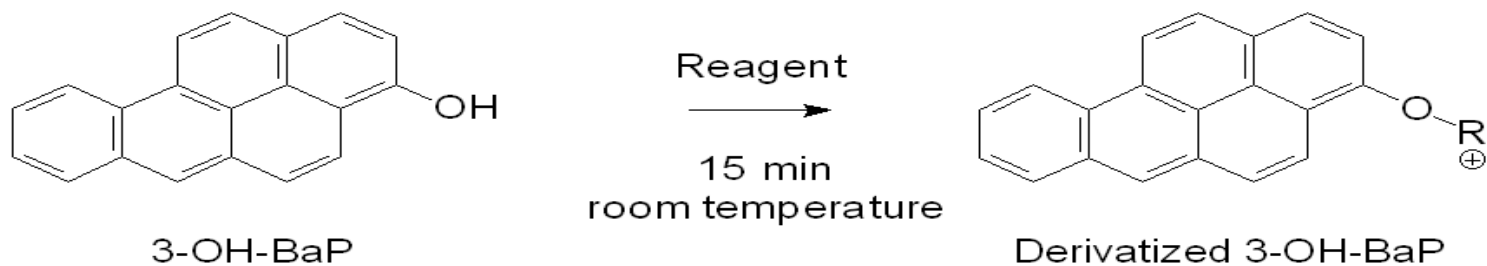


# Sub-pg/mL Method Example



3-Hydroxybenzo[a]pyrene  
 $C_{20}H_{12}O$   
268.31 Da

# Derivatization of 3-OH-BaP

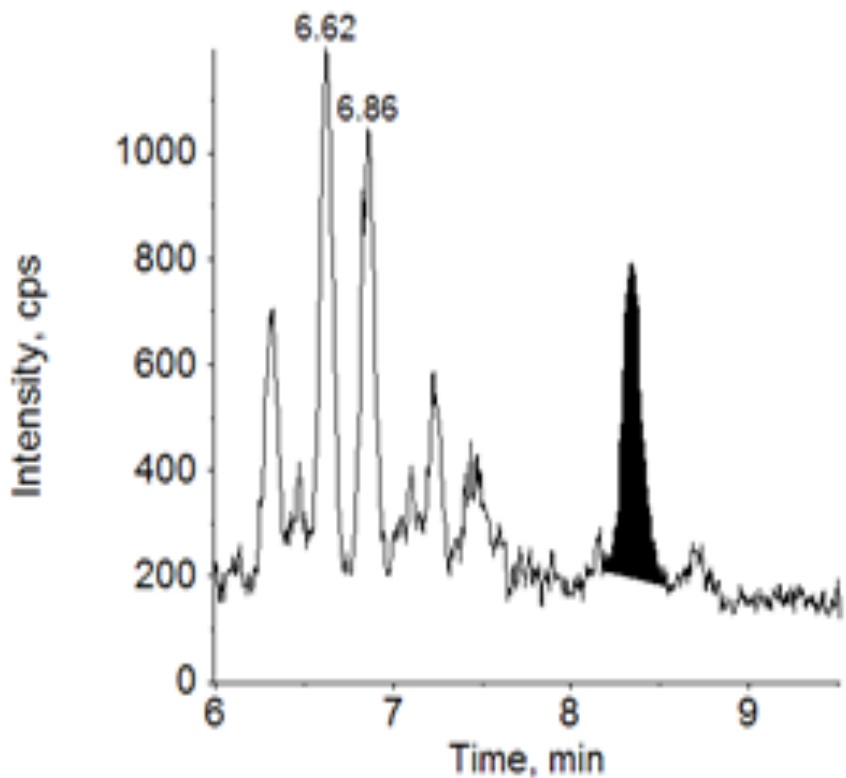


Derivatization added a new functional group which is easily ionized.

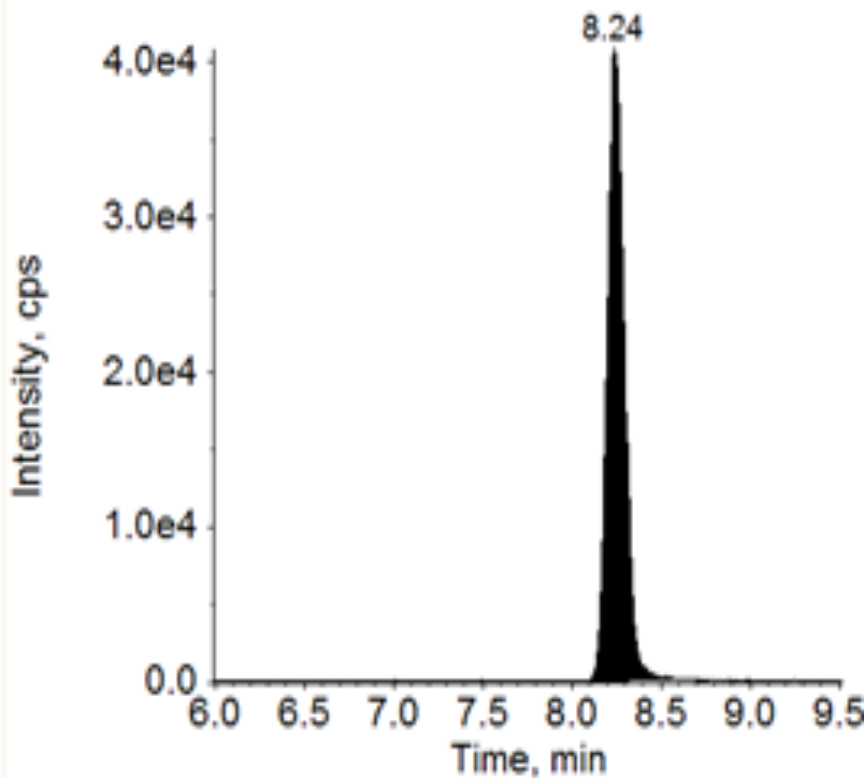
Improved fragmentation- MS/MS efficiency in positive ion mode.

# 3-OH BaP LLOQ S/N= 23 (50 fg/mL in dog urine )

107 010 AA04030-01 STD B 1 1 - 3-Hydroxyb...  
Area: 4553.530 counts Height: 5.95e+002 c...

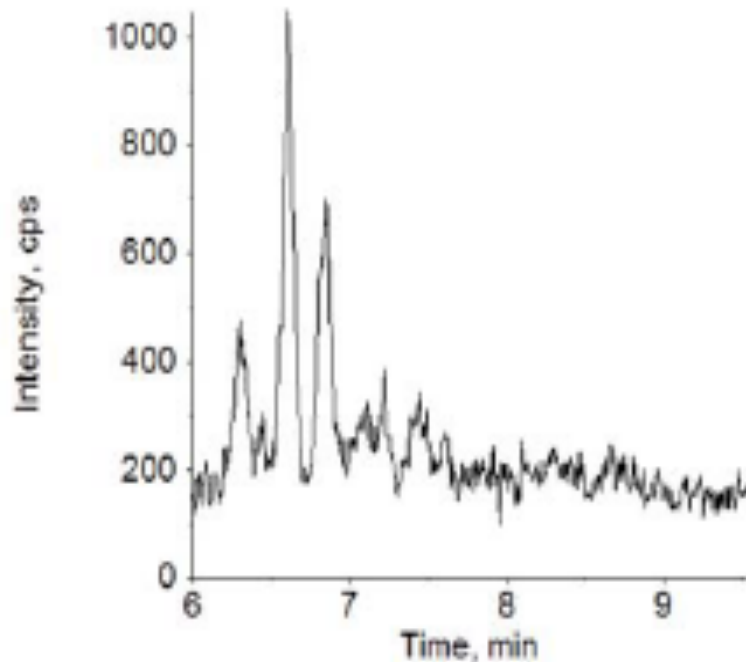


107 010 AA04030-01 STD B 1 1 - d5-3-OH-B...  
Area: 270496.700 counts Height: 4.07e+00...

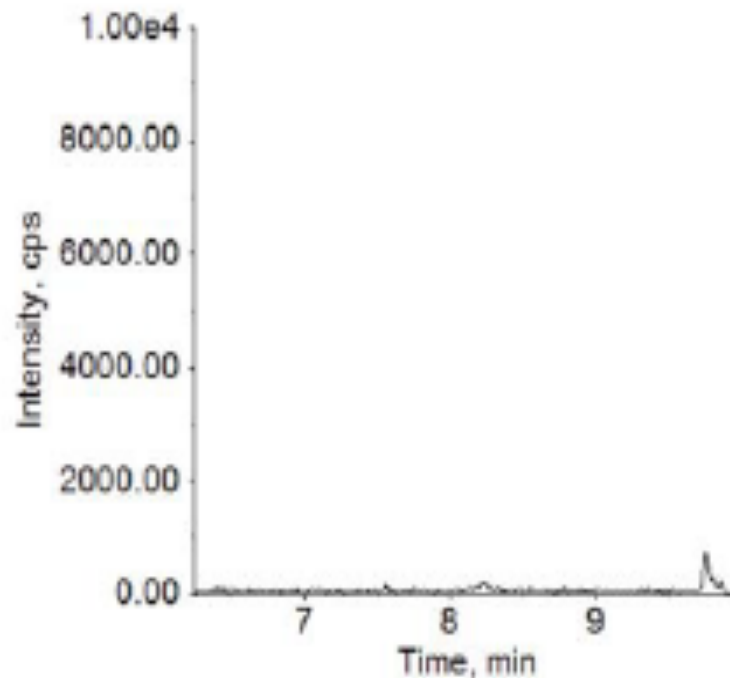


# 3-OH BaP Extracted Blank (in dog urine)

107 008 AA04030-01 Control Blank 1 - 3-Hyd...  
(peak not found)



107 008 AA04030-01 Control Blank 1 - d5-3-...  
(peak not found)

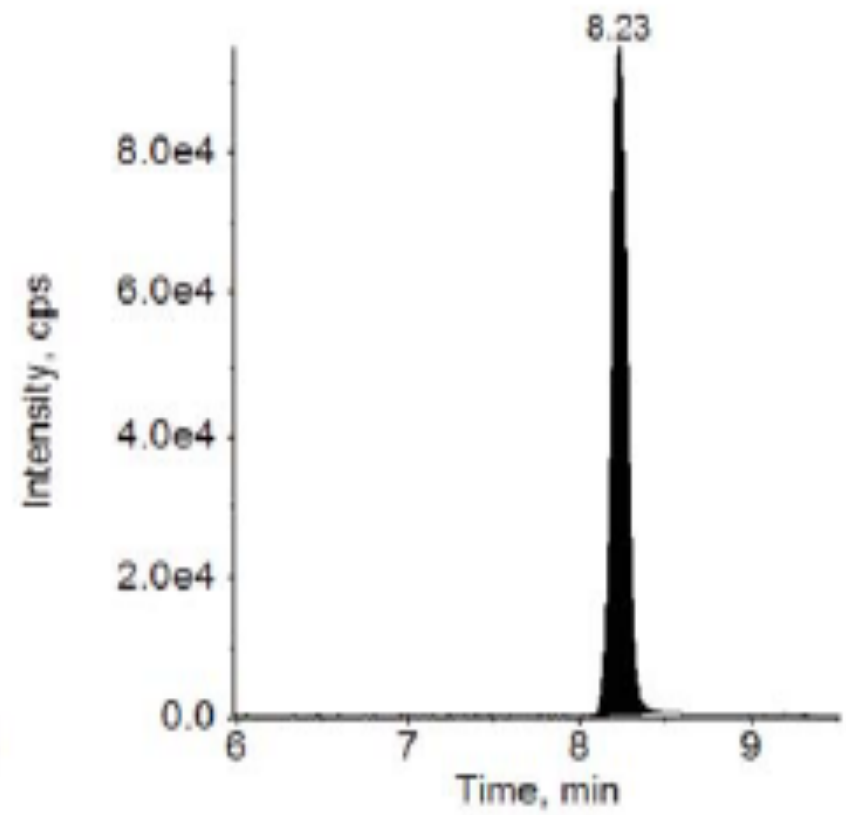
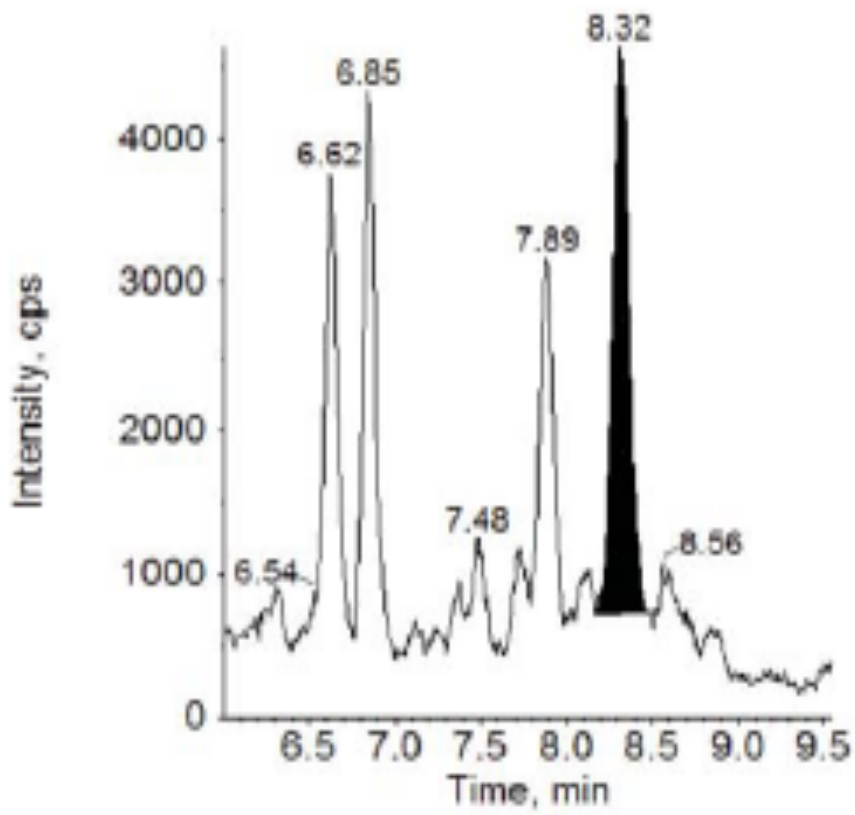




# 3-OH BaP Extracted Human Urine Sample Approximately 150 fg/mL

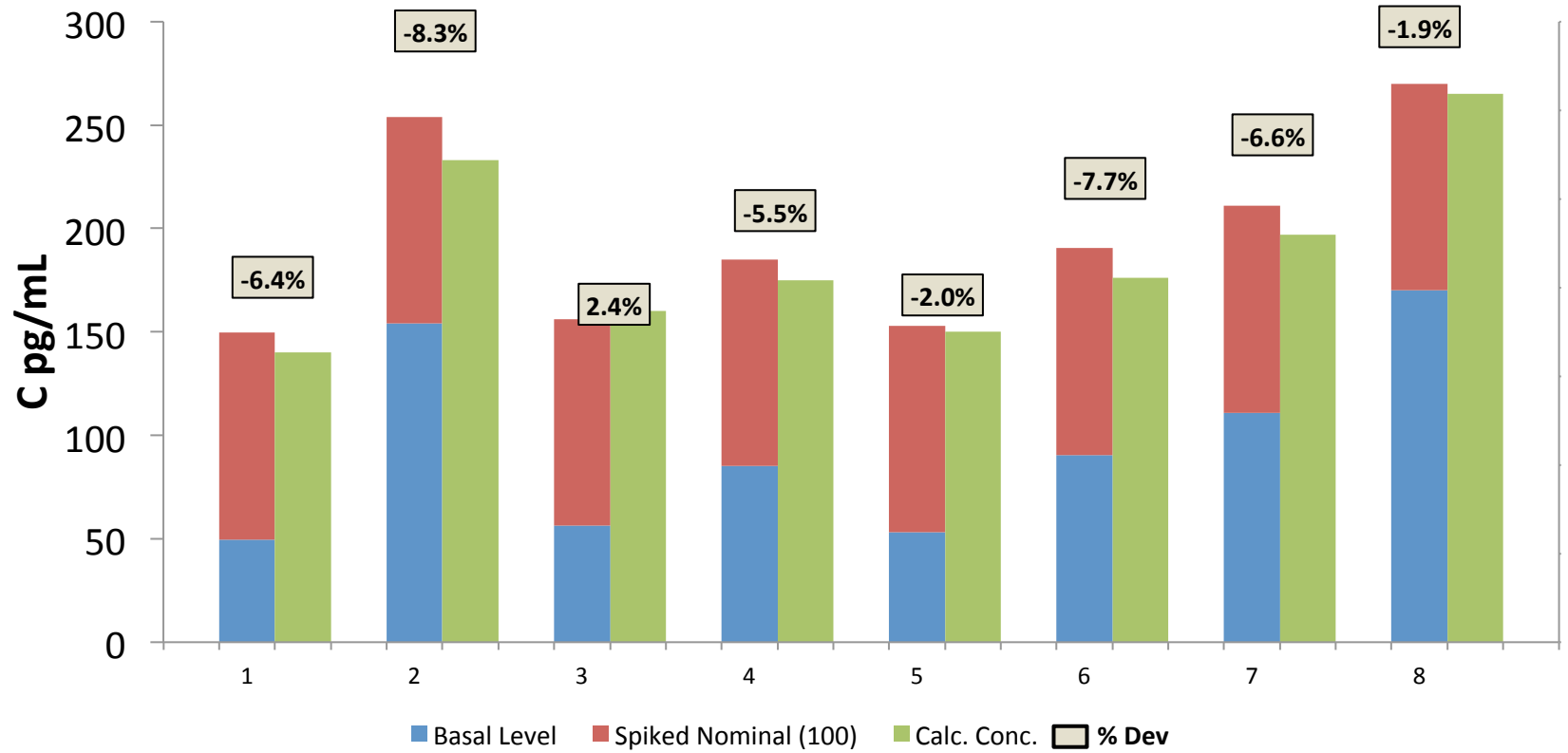
120 045 AA04030-01 TEST A 1 - 3-Hydro...  
Area: 25435.508 counts Height: 3.93e+...

120 045 AA04030-01 TEST A 1 - d5-3-OH-...  
Area: 571089.443 counts Height: 9.45e+...



# Accuracy of Matrix Effect Data in Human Urine

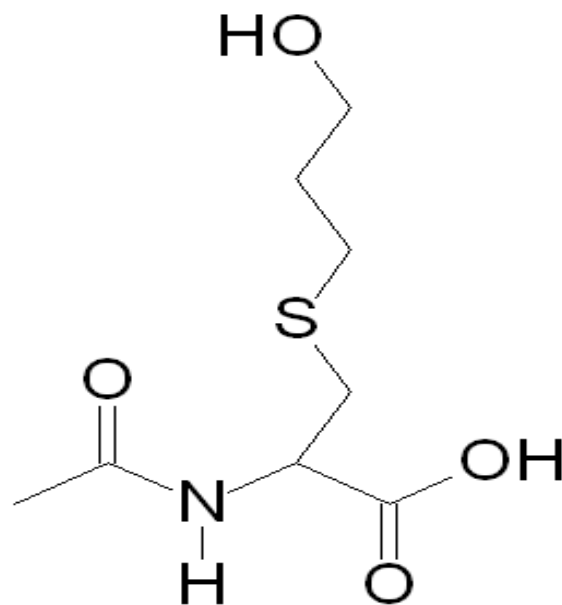
### Selectivity and Matrix Effect Data in Ten Lots of Human Urine for 3-OH-BaP



# Summary for 3-OH-BaP Method

- An ultra-sensitive method for 3-OH-B[a]P with LLOQ of 50 fg/mL was developed and validated.
- Robustness was demonstrated by a batch passing rate over 98%.

# Structures of 3-HPMA

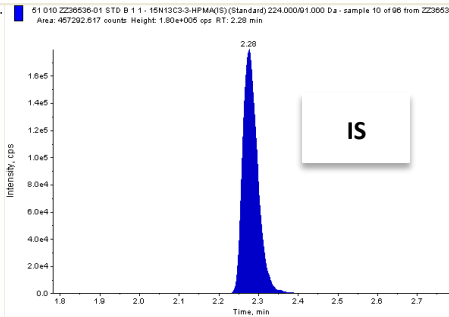
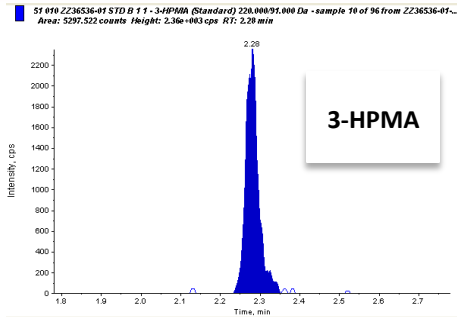


3-HPMA

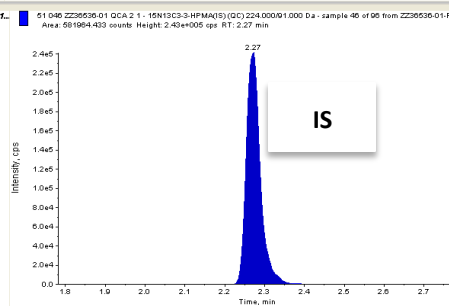
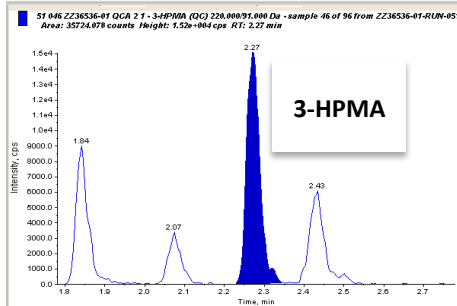
N-Acetyl-S-(3-hydroxypropyl)cysteine

221.28 Da

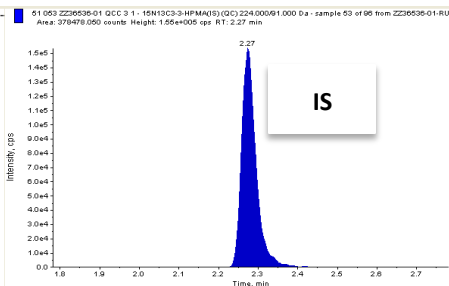
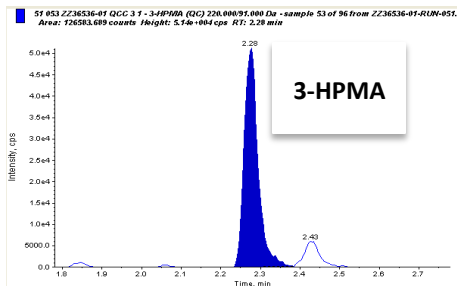
# 3-HPMA RP-UHPLC Chromatography Using the Stable labeled Internal standard



LLOQ in Artificial Urine

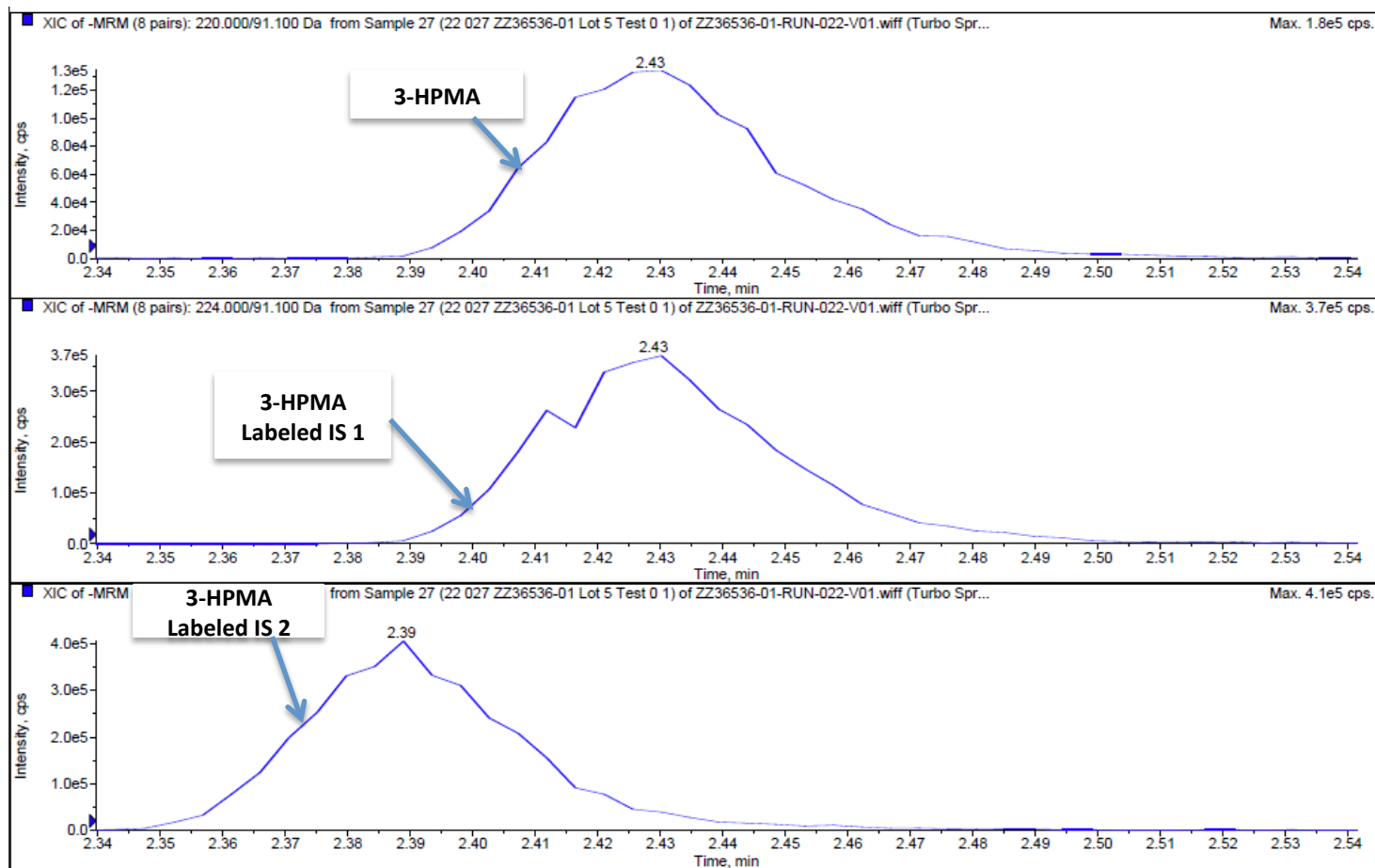


Low QC in Human Urine



High QC in Human Urine

# 3-HPMA UPLC RP chromatography using 2 stable labeled internal standards



# Matrix Effect Test in Human Urine Comparing the Performance of Two Different Labeled Internal Standards

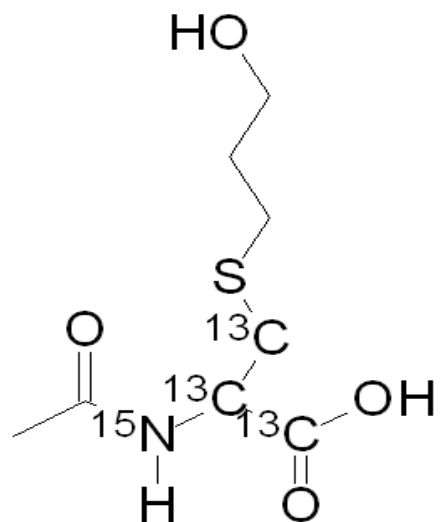
A

Lot#	Basal Level (ng/mL)	Low Spiked Nominal Conc. (ng/mL)	Expected Conc. (ng/mL)	Calc. Conc. (ng/mL)	% Dev.	High Spiked Nominal Conc. (ng/mL)	Expected Conc. (ng/mL)	Calc. Conc. (ng/mL)	% Dev.
1	813	20.0	833	768	-7.8	3750	4563	4660	+2.1
2	1160	20.0	1180	1150	-2.5	3750	4910	4840	-1.4
3	31.1	20.0	51.1	53.1	+3.9	3750	3781	3900	+3.1
4	225	20.0	245	256	+4.5	3750	3975	4010	+0.9
5	22.1	20.0	42.1	39.3	-6.7	3750	3772	3950	+4.7
6	1320	20.0	1340	1280	-4.5	3750	5070	5160	+1.8
7	1960	20.0	1980	1850	-6.6	3750	5710	5660	-0.9
8	440	20.0	460	446	-3.0	3751	4191	4130	-1.5

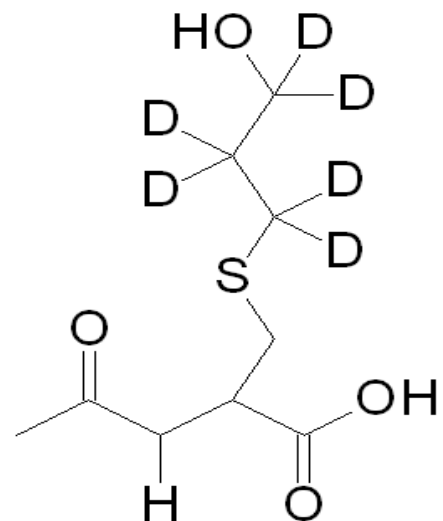
B

Lot#	Basal Level (ng/mL)	Low Spiked Nominal Conc. (ng/mL)	Expected Conc. (ng/mL)	Calc. Conc. (ng/mL)	% Dev.	High Spiked Nominal Conc. (ng/mL)	Expected Conc. (ng/mL)	Calc. Conc. (ng/mL)	% Dev.
1	795	20.0	815	828	+1.6	3750	4545	4080	-10.2
2	902	20.0	922	860	-6.7	3750	4652	3500	<b>-24.8</b>
3	30.7	20.0	50.7	49.1	-3.2	3750	3781	3390	-10.3
4	227	20.0	247	268	+8.5	3750	3977	4140	+4.1
5	21.3	20.0	41.3	38.4	-7.0	3750	3771	3790	+0.5
6	1230	20.0	1250	1110	-11.2	3750	4980	4260	<b>-14.5</b>
7	2330	20.0	2350	2300	-2.1	3750	6080	6970	<b>+14.6</b>
8	536	20.0	556	572	+2.9	3750	4286	5270	<b>+23.0</b>

# Structures of 3-HPMA Internal Standards



$[^{15}\text{N}, ^{13}\text{C}_3]$ -3-HPMA  
225.25 Da



$[^2\text{H}_6]$ -3-HPMA  
226.32 Da

$d_6$ -3HPMA failed to accurately quantify 3-HPMA matrix effect test in human urine when  $[^{15}\text{N}^{13}\text{C}_3]$ 3-HPMA reliably tracked the analyte.



# 3-HPMA Method Results

- A LC-MS/MS method for the simultaneous analysis of CEMA, 3-HPMA and HBMA with improved selectivity has been developed and validated.

# Conclusion

- Two dimensional chromatography provided better selectivity for the assay of NNN and MHBMA.
- Derivatization provided an excellent solution to enhancing sensitivity to reach sub pg/mL levels 3-OH-BaP.
- $^{13}\text{C}$  and  $^{15}\text{N}$  labeled internal standards which coelute with the analytes provide the most robust bioanalytical methods for 3-HPMA.

# Acknowledgments

- Alan Dzerk
- Christine Kafonek
- Kirk Newland
- Patrick Miller
- Veni Lapko
- Paul Brown



# **The Science Required For Successful Bioanalytical Methods**

**By: Ridha Nachi**

**TSRC, 2012**