

# Assuring the Quality of MS/MS Dried Blood Spot Newborn Screening Testing

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# National Impact of CDC's Newborn Screening Quality Assurance Program (NSQAP)

- ❑ Sole provider of comprehensive quality assurance services for screening labs
  - Essential for evaluation of method performance
- ❑ Over 850,000 dried-blood spots (DBS) produced each year
  - 3 challenges of 5 blind-coded samples/year
  - Proactive follow-up of false negative results
- ❑ 100% of US states covered by program
  - Allows for accreditation of screening labs
  - Provide summary reports and feedback for all participating labs



# Available MS/MS-Detectable Markers

- ❑ Alanine
- ❑ Citrulline
- ❑ Phenylalanine
- ❑ Leucine
- ❑ Valine
- ❑ Methionine
- ❑ Arginine
- ❑ Tyrosine
- ❑ Succinylacetone (tyrosinemia type I)
- ❑ Free carnitine (C0)
- ❑ Acetylcarnitine (C2)
- ❑ Propionylcarnitine (C3)
- ❑ Malonylcarnitine (C3DC)
- ❑ Isobutyrylcarnitine (C4)
- ❑ 3-Hydroxyisobutyrylcarnitine (C4OH)
- ❑ Isovalerylcarnitine (C5)
- ❑ Glutaryl carnitine (C5DC)
- ❑ Tiglylcarnitine (C5:1)
- ❑ 3-Hydroxyisovalerylcarnitine (C5OH)
- ❑ Hexanoylcarnitine (C6)
- ❑ Octanoylcarnitine (C8)
- ❑ Decanoylcarnitine (C10)
- ❑ Decenoylcarnitine (C10:1)
- ❑ Decadienoylcarnitine (C10:2)
- ❑ Dodecanoylcarnitine (C12)
- ❑ Myristoylcarnitine (C14)
- ❑ Tetradecenoylcarnitine (C14:1)
- ❑ 3-Hydroxypalmitoylcarnitine (C16OH)
- ❑ Palmitoylcarnitine (C16)
- ❑ Stearoylcarnitine (C18)
- ❑ Oleoylcarnitine (C18:1)
- ❑ Androstenedione/cortisol/11-deoxy/21-deoxy
- ❑ ABG, ASM, GAA, GALC, GLA, IDUA



## Comparison of amino acids and acylcarnitines assay methods used in newborn screening assays by tandem mass spectrometry

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- ❑ Minor differences found for most analytes
- ❑ Arg, Cit can differ also depending on method
- ❑ C3DC is most different
- ❑ NSQAP Data Examples

### 4. Discussion

Overall differences between amino acid and acylcarnitine quantitative values resulting from derivatized and underivatized techniques were <15% for the majority of the amino acids and acylcarnitines detected by both methods. However, the most striking difference between the MS/MS analysis of acylcarnitines by butyl esterification techniques as compared to underivatized forms is for dicarboxylic species such as C5DC. These differences can be explained, in part, based on the results from a previous study of C3DC [1]. Laboratories report concentrations based on ion ratios of the unlabeled dicarboxylic acylcarnitine to an internal standard. That internal standard is

Figure 15. Bias Plot of Phenylalanine Values by Method  
Quarter 3, Specimen 3  
Expected Value (EV) 295.25  $\mu\text{mol/L}$  whole blood

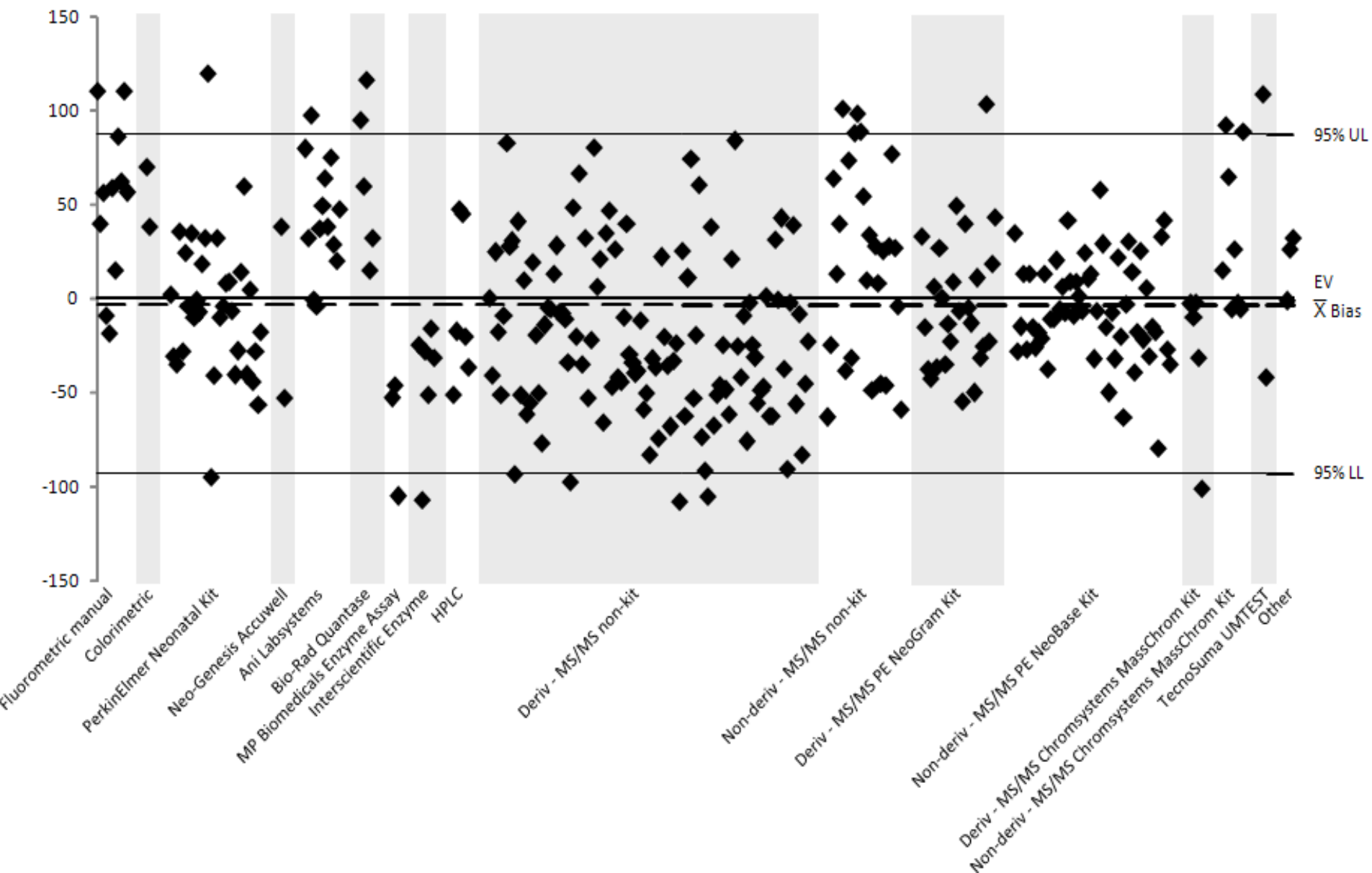


Figure 23. Bias Plot of Free Carnitine (C0(L)) Values by Method  
Quarter 1, Specimen 1  
Expected Value (EV) 5.66  $\mu\text{mol/L}$  whole blood

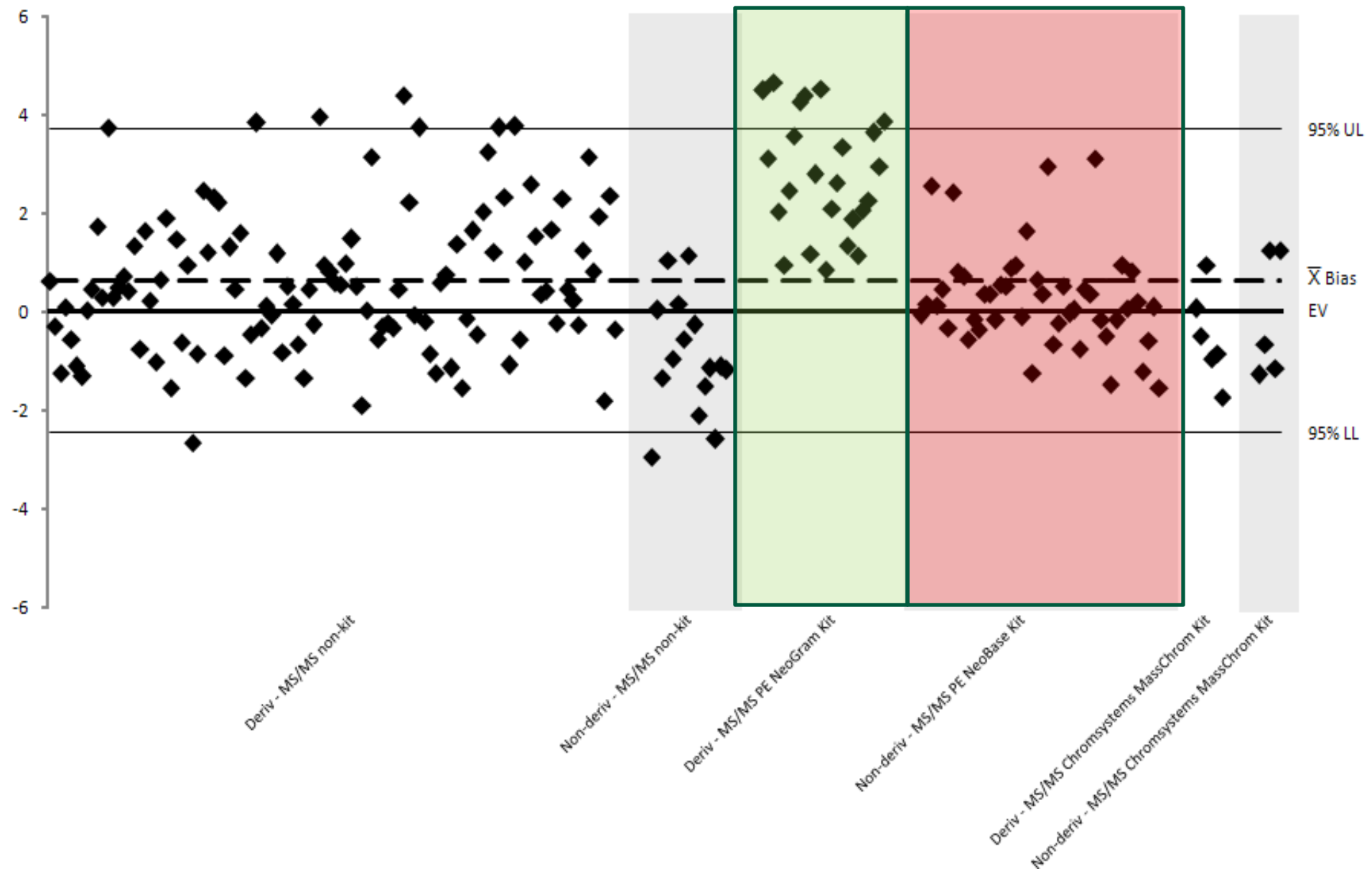
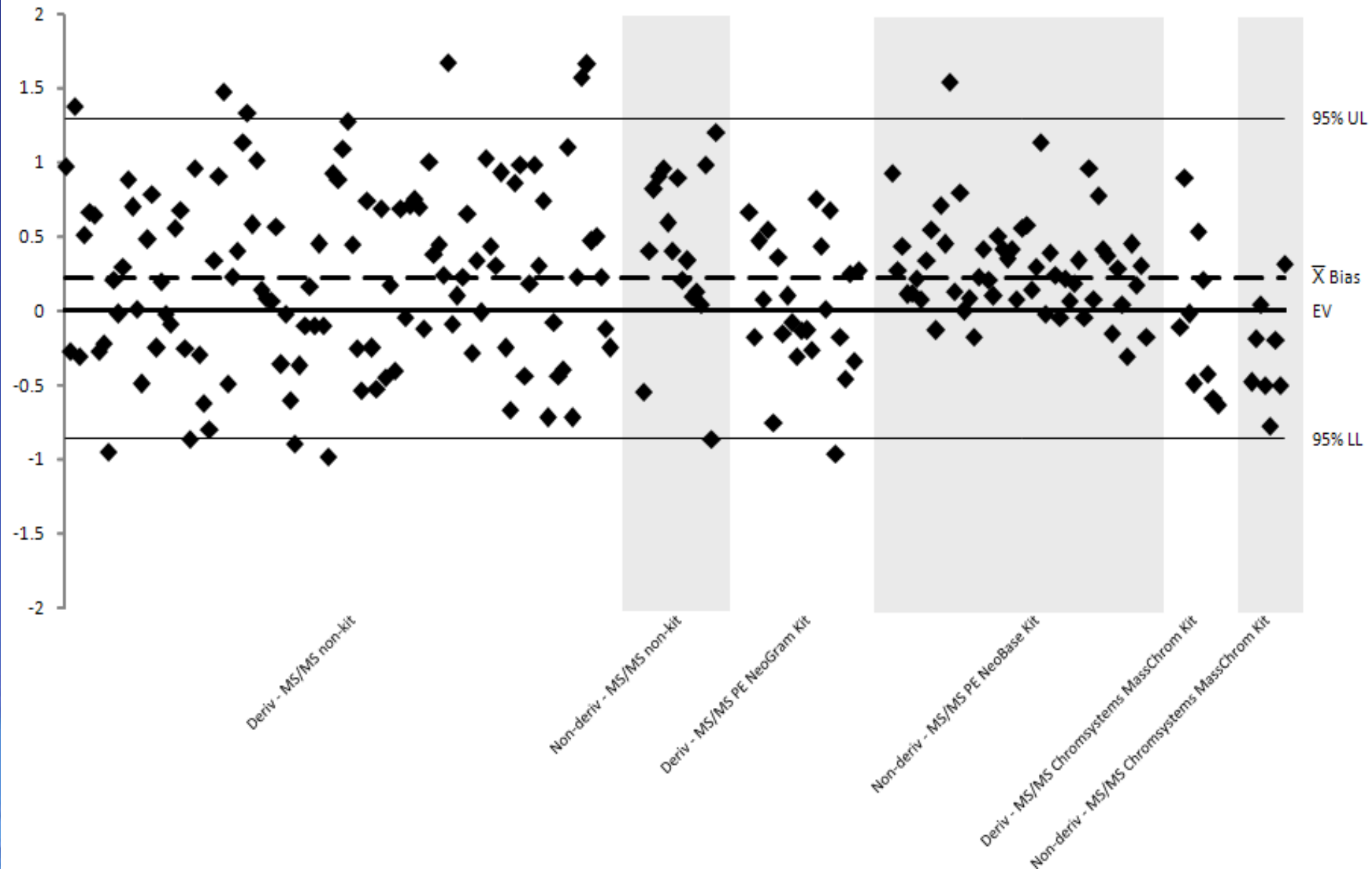


Figure 33. Bias Plot of Octanoylcarnitine (C8) Values by Method  
Quarter 3, Specimen 5  
Expected Value (EV) 3.10  $\mu\text{mol/L}$  whole blood







# Malonylcarnitine (C3DC): The beginning

- Dicarboxylic acylcarnitine biomarker used to screen for malonic acidemia (MAL)
  - IEM caused by congenital deficiency of malonyl-CoA decarboxylase
  - (2<sup>nd</sup> target panel)
  
- Introduced into NSQAP PT panels in 2008

**Table 2**  
Newborn screening panel: core panel and secondary targets

MS/MS				
Acylcarnitines		Amino acids		
9 OA	5 FAO	6 AA	3 Hb Pathies	6 Others
<b>CORE PANEL</b>				
IVA	MCAD	PKU	Hb SS*	CH
GA I	VLCAD	MSUD	Hb S/βTh*	BIOT
HMG	LCHAD	HCY*	Hb S/C*	CAH*
MCD	TFP	CIT		GALT
MUT*	CUD	ASA		HEAR
3MCC*		TYR I*		CF
Cbl A,B*				
PROP				
BKT				
<b>SECONDARY TARGETS</b>				
6 OA	8 FAO	8 AA	1 Hb Pathies	2 Others
Cbl C,D*	SCAD	HYPER-PHE	Var Hb*	GALK*
<b>MAL</b>	GA2	TYR II		GALE
IBG	M/SCHAD	BIOPT (BS)		
2M3HBA	MCKAT	ARG		
2MBG	CPT II	TYR III		
3MGA	CACT	BIOPT (REG)		
	CPT IA	MET		
	DE RED	CIT II		

NOTE: Codes are as follows: OA, disorders of organic acid metabolism; FAO, disorders of fatty acid metabolism; AA, disorders of amino acid metabolism; Hb Pathies, hemoglobinopathies.  
\* Identifies conditions for which specific discussions of unique issues are found in the main report.

# And then there were issues with C3DC analysis...

## □ Non-derivatized assay

- Kit-, non-kit-based
- Lower semi-quantitative results

## □ Hydroxybutyrylcarnitine (C4OH)

- Introduced into NSQAP PT panels in 2010
- Isobaric interference
  - m/z 248

## □ PT misses ensued

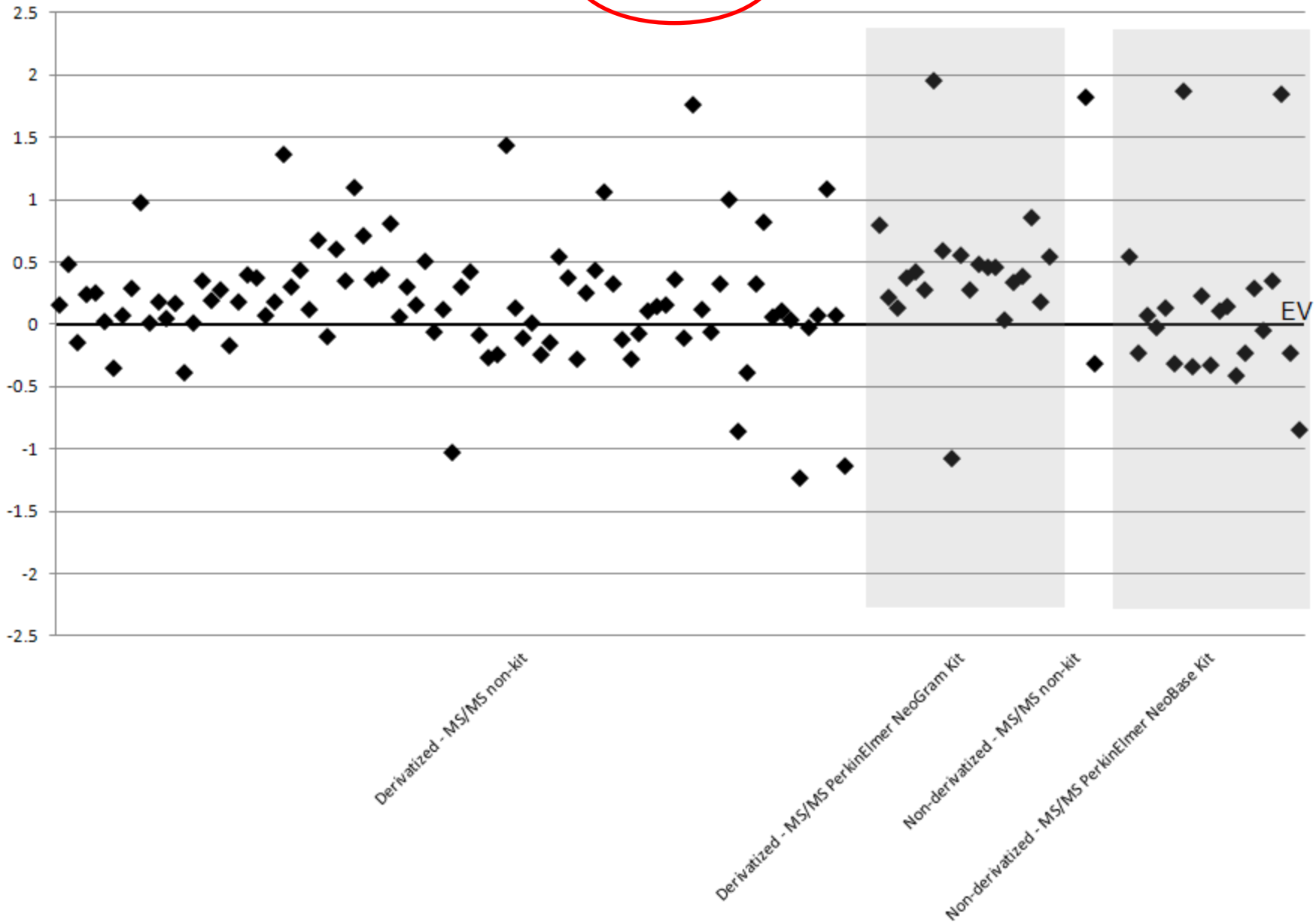
- Increased corrective action reports
- General feeling: WTH?



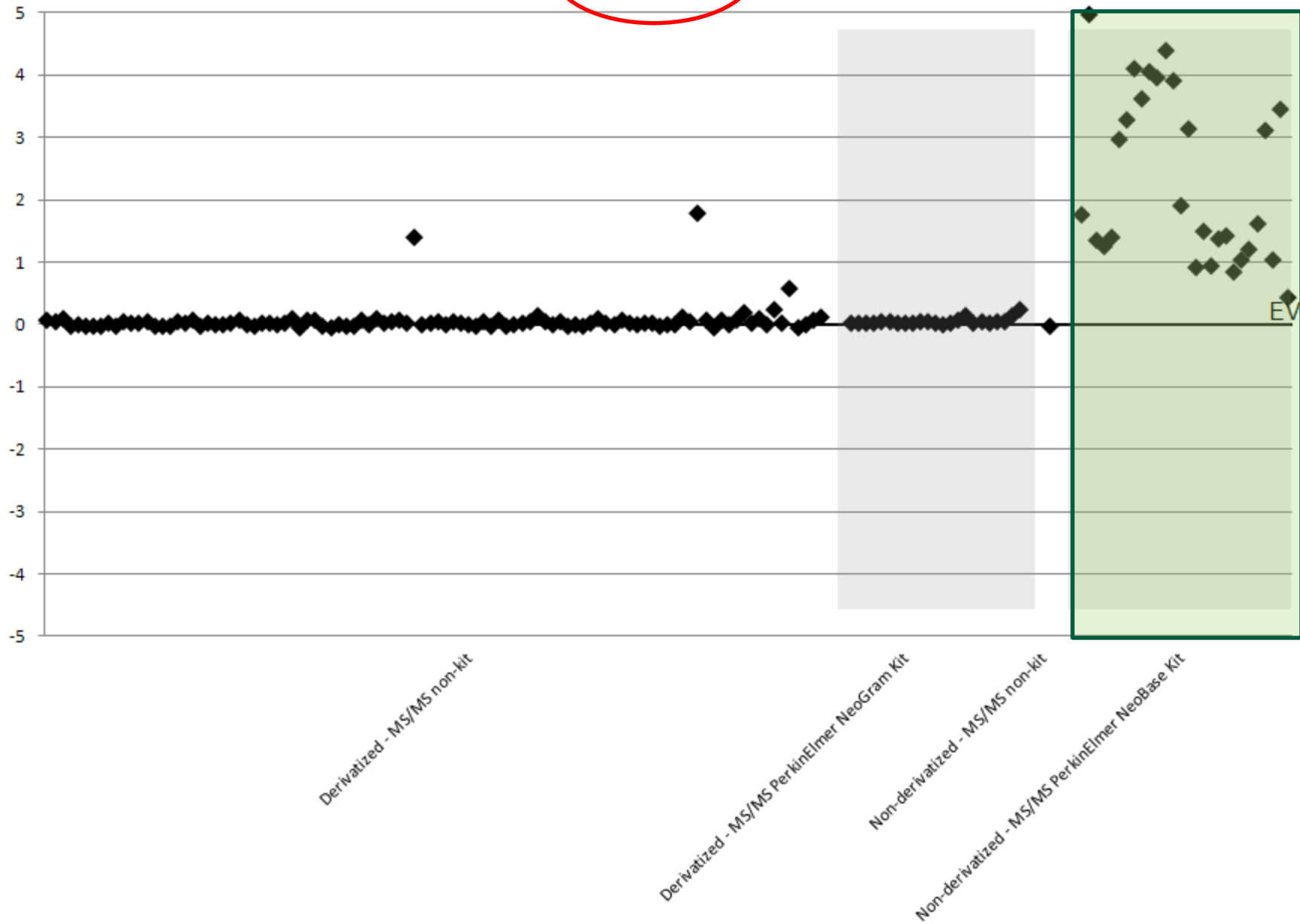
# Bland Altman Plot: Hydroxybutyrylcarnitine (C4OH)

Quarter 3, Specimen 3161

Expected Value (EV) 1.33  $\mu\text{mol/L}$  whole blood



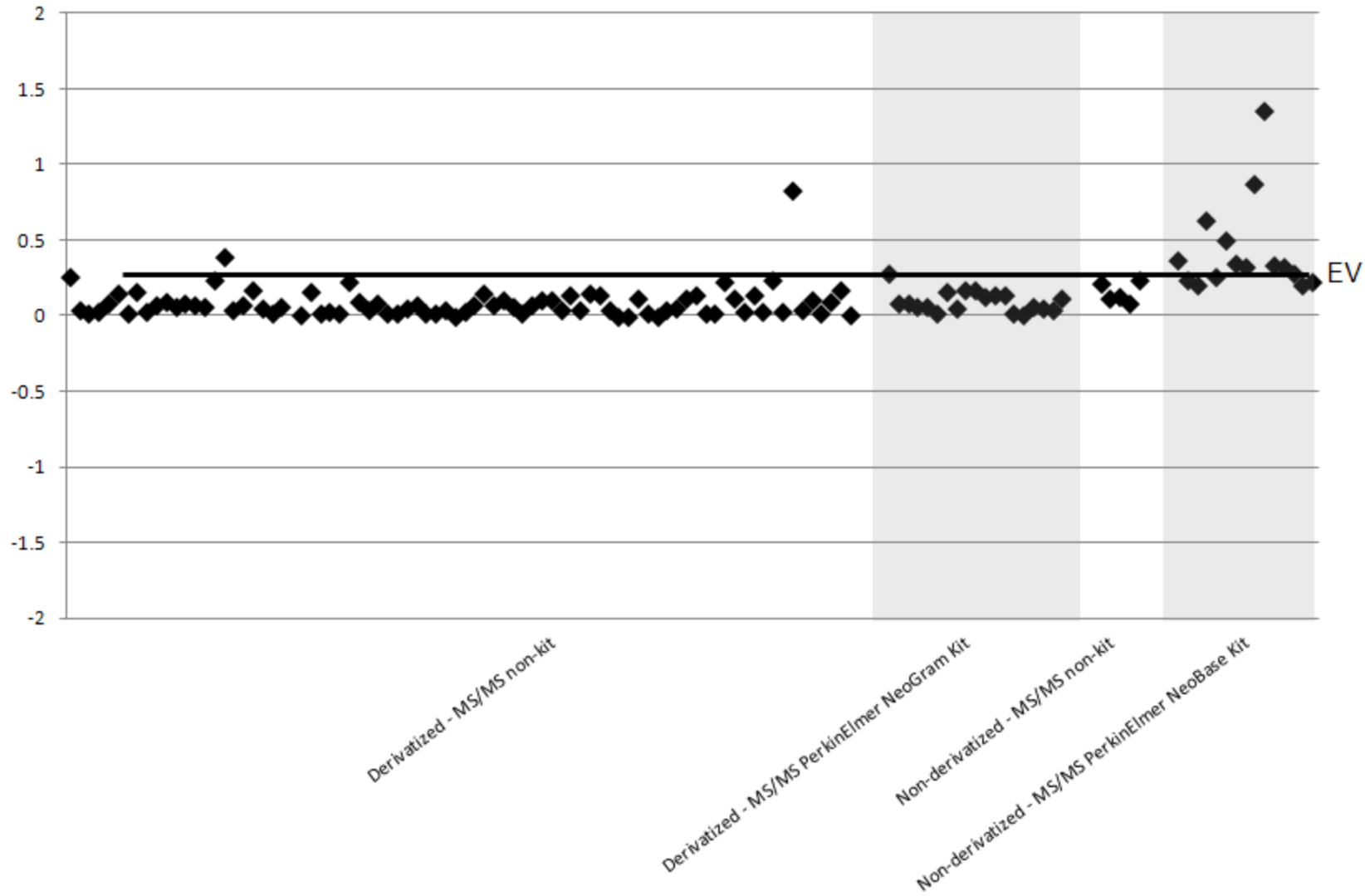
Bland Altman Plot: Malonylcarnitine (C3DC)  
Quarter 3, Specimen 3161  
Expected Value (EV) 0.07  $\mu\text{mol/L}$  whole blood



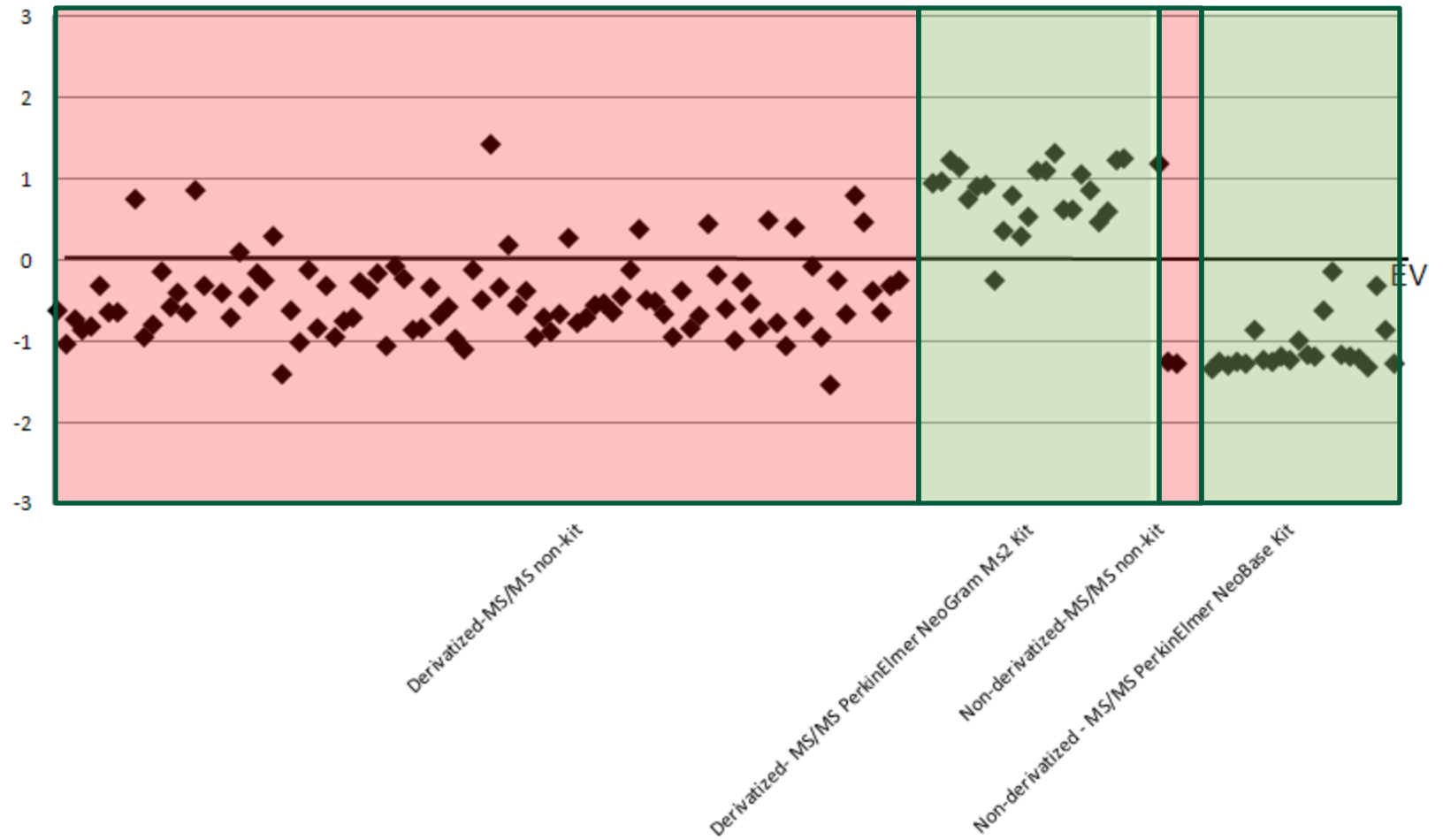
# Bland Altman Plot: Hydroxybutyrylcarnitine (C4OH)

Quarter 1, Specimen 1163

Expected Value (EV) = 0.07  $\mu\text{mol/L}$  whole blood



Bland Altman Plot: Malonylcarnitine (C3DC)  
Quarter 1, Specimen 1163  
Expected Value (EV) = 1.57  $\mu\text{mol/L}$  whole blood



**Ionization Efficiency**

# MS/MS Performance Metrics 2006 - 2011

Selected Domestic False Positive Rates (%) for 2006-2011

Disorder/Analyte	Year					
	2006	2007	2008	2009	2010	2011
Phenylketonuria (Phe)	0.2	0.5	0.7	1.8	0.1	0.3
Maple Syrup Urine Disease (Leu)	0.6	3.1	0.7	0.6	0.1	0.2
Tyrosinemia I, II, III (Tyr)	0.0	0.3	0.0	0.3	0.3	0.0
Maple Syrup Urine Disease (Val)	2.2	1.7	0.4	0.7	0.1	1.6
Citrullinemia (Cit)	1.1	0.0	0.2	0.4	0.4	0.3
C3 Screen	0.0	0.1	0.3	0.8	0.3	0.0
C3DC Screen	N/A		0.8	1.6	2.9	0.3
C5 Screen	0.0	0.0	0.9	1.0	0.1	0.0
C5DC Screen	0.0	0.0	0.0	1.0	0.0	0.0
C8 Screen	0.1	0.3	0.0	0.8	0.1	0.3
C16 Screen	0.0	0.1	0.0	0.4	0.2	0.0

# MS/MS Performance Metrics 2006 - 2011

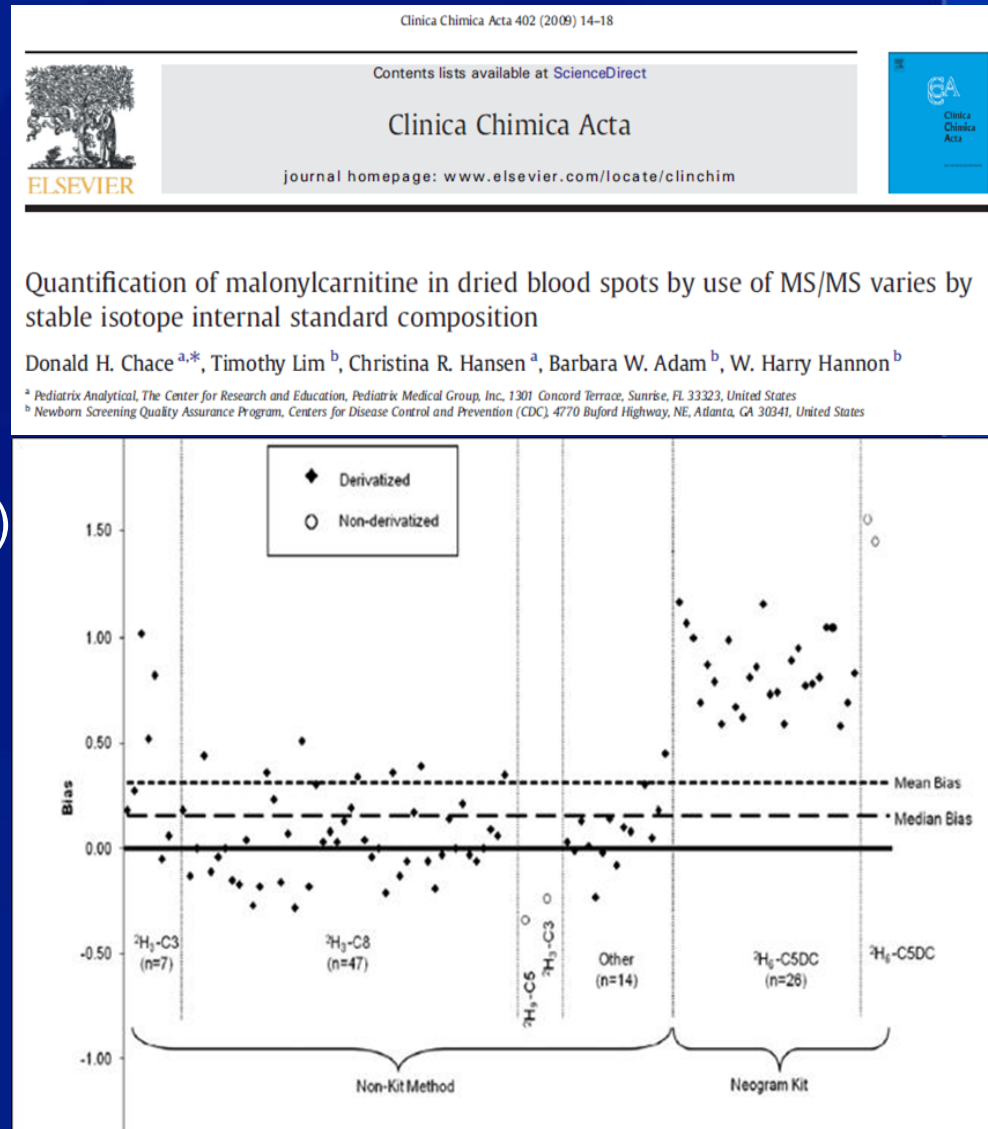
Selected Domestic False Negative Rates (%) for 2006-2011

Disorder/Analyte	Year					
	2006	2007	2008	2009	2010	2011
Phenylketonuria (Phe)	0.6	0.0	1.1	0.5	0.8	0.0
Maple Syrup Urine Disease (Leu)	0.0	0.0	0.0	1.1	0.5	0.0
Tyrosinemia I, II, III (Tyr)	1.6	0.7	3.3	1.0	1.5	0.0
Maple Syrup Urine Disease (Val)	0.0	0.0	0.0	0.9	1.1	0.0
Citrullinemia (Cit)	0.0	0.0	0.0	1.7	0.5	0.0
C3 Screen	1.9	0.0	0.0	2.1	0.7	0.0
C3DC Screen	N/A		0.0	4.0	19.4	11.7
C5 Screen	0.8	0.0	0.0	4.0	0.5	0.0
C5DC Screen	3.7	0.0	0.0	1.7	1.0	0.6
C8 Screen	0.6	0.0	0.0	1.2	0.7	0.0
C16 Screen	0.6	0.0	0.0	8.9	1.0	2.0



# What to do?

- ❑ **Derivatized assay can resolve C3DC and C4OH!**
  - C3DC, C5DC analysis enhanced by derivatization
  - Choice of IS (Chace et al 2009)
  - If unable to derivatize, establish ratios, work with other labs
  - Follow-up procedures for correct screening classification (i.e., cutoffs)



# NSQAP adapts to ensure high-quality screening

## □ PT Testing

- NSQAP new category: C3DC + C4OH
- Allows for reduced corrective action reports
- No double-dipping!
- On-line reporting category: live in January 2012

## □ QC Materials

- Two characterization sheets for AA, AC QC materials
- No changes to reporting scheme

# Summary

- ❑ **Newborn screening by tandem mass spectrometry is a successful public health program**
  - >95% of newborns screened in US
  - 42/56 RUSP disorders are MS/MS-detectable
- ❑ **Many challenges remain for DER-UND screening**
  - Understanding assay and metabolite limitations is key
  - Establish proper procedures to eliminate false positives and negatives
- ❑ **NSQAP is a comprehensive resource for laboratory services**
  - New PT reporting reflects current practices in the field

## For More Information

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NSQAP Web Site:

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