

#### NATTS TAD Revision 3

**Method Detection Limits** 

Method Update Rule vs 40 CFR Part 136



# **Method Detection Limits**

- Agenda:
  - Background of detection limits
  - MDLs as they relate to the NATTS network
  - MDL Method Update Rule
  - Determining MDLs via Method Update Rule
  - Analysis of reported AQS data to examine impact of MUR





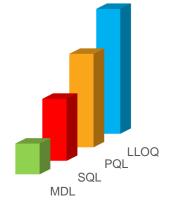
- The <u>method detection limit</u> (MDL) is defined as the minimum concentration of a substance that can be measured and reported with 99% confidence that the analyte concentration is greater than zero and is determined from analysis of a sample in a given matrix containing the analyte.
- Initially designated for water analyses, but adapted for many other matrices.





# **Detection Threshold Terms**

- Common terms related to the detection threshold
  - Method Detection Limit (MDL) defined by 40 CFR Part 136 App B
  - Limit of Detection (LOD) TNI term for detection, not quantitation
  - Sample Quantitation Limit (SQL) typically ~3x the MDL
  - Practical Quantitation Limit (PQL) typically 5x the MDL
  - Lower Limit of Quantitation (LLOQ) typically the lowest calibration standard level
  - Method Reporting Limit (MRL) lowest quantifiable concentration, often set at the LLOQ





# **Method Detection Limits**

- MDLs are controversial
- You may have heard (or said yourself):
  - "MDLs are flawed because they are a theoretical value."
  - "MDLs don't reflect real-world conditions."
  - "MDLs don't take into account the sample media background."
  - "We don't trust values around the MDL."





# **NATTS Network MDLs**

- MDLs are necessary to properly assess risks, particularly at the ambient concentrations measured by the NATTS network
- Concentration data must be qualified according to relationship with MDL and SQL so data users can assign proper confidence
- Consistent determination of analyte MDLs across all NATTS laboratories ensures results are comparable
- MDLs aren't going away anytime soon



#### **Method Detection Limits**

 40 CFR Part 136 App B prescribes preparation and analysis of a minimum of 7 spikes in matrix. The MDL is calculated by multiplying the standard deviation of the measured concentrations by the appropriate student's T.

$$MDL = s \cdot T$$

• Compare the determined MDL to the nominal spiked value.

MDL < spike value < 10x MDL



# NATTS MDLs – TSAs



- TSAs of NATTS network laboratories found that very few laboratories determined MDLs correctly per 40 CFR Part 136 Appendix B or were not determining them annually.
  - Analyzed a standard directly without matrix 7 times
  - Analyzed one or two samples annually to "verify" the MDL
  - Prepared one spiked sample and analyzed this sample 7 times
  - Prepared 7 spiked samples not including all portions of the matrix
  - Did not ensure that the nominal spiked value was 1- to 10-fold the determined MDL

# MDLs – Method Update Rule

- Draft Revision 3 of the NATTS TAD adopts the Method Update Rule
- Why adopt the Method Update Rule...?
  - MDLs determined by 40 CFR Part 136 Appendix B do not take into account the sample collection media.
  - TO-11A permits up to 0.150 μg/cartridge, or 0.104 μg/m<sup>3</sup> assuming 1.44 m<sup>3</sup> collected volume. This exceeds the MDL MQO of 0.08 μg/m<sup>3</sup>.
  - An analysis of AQS data and laboratory reported MDL data indicate that little to no additional data would be qualified as < MDL following adoption of the MUR.</li>





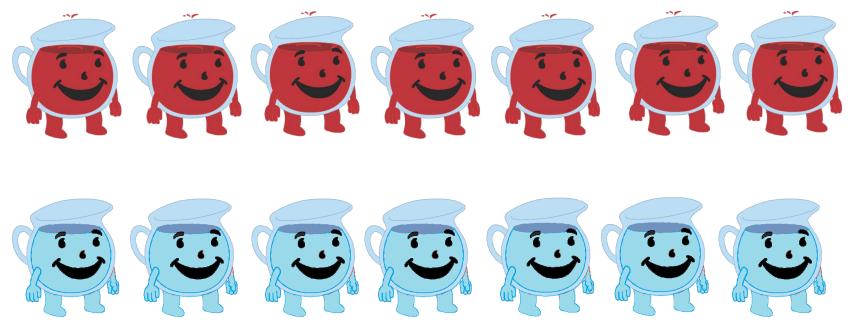
# MDLs – Method Update Rule

- Improvements to 40 CFR Part 136 Appendix B:
  - Incorporates the background attributable to the sample collection media
  - More closely approximates real-world conditions
    - Incorporates temporal variability requires minimally three separate batches for sample preparation and analysis over minimally three separate dates
- Still requires sample preparation in a laboratory in a clean matrix
- Does not address that the determined MDL is theoretical



# MDL – Method Update Rule

• Prepare a minimum of 7 spikes and 7 blanks over the course of three or more different preparation batches





# MDL – Method Update Rule

• Determine the MDL as per 40 CFR Part 136 Appendix B

$$MDL_{sp} = S_{sp} \cdot T$$

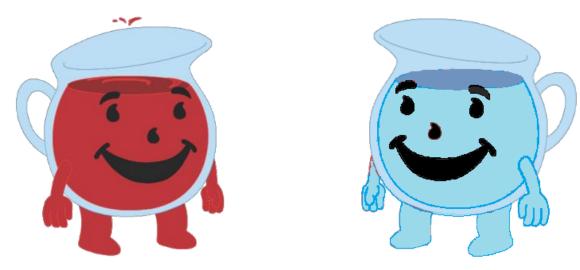
- Determine an additional MDL based on the analysis of the blanks
  - Calculate the average  $(\bar{x}_b)$  and standard deviation  $(s_b)$  of the blanks.
  - Multiply the blank standard deviation (s<sub>b</sub>) by the appropriate student's T value and add this to the average blank value ( $\bar{x}_b$ ).

$$MDL_b = (s_b \cdot T) + \overline{x}_b$$



#### **MDL- Method Update Rule**

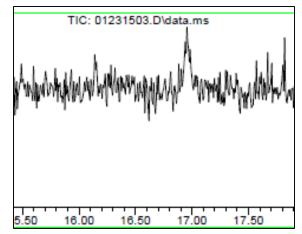
- Compare MDL<sub>sp</sub> and MDL<sub>b</sub>
- Whichever is higher is reported as the laboratory MDL





#### **MDLs – Determining Spike Levels**

- Choose a spiking level by considering:
  - Previously acceptable MDL studies and related experience
  - Concentration where qualitative identification criteria are lost
  - Concentration at which the signal to noise ratio is ~3 to 5-fold
  - Analysis of a suite of blank samples calculate the standard deviation and multiply by 3





- MDLs (mean or median) were polled from laboratories participating in the NATTS PT for metals, VOCs, PAHs, and carbonyls.
- These "typical" laboratory MDLs were all below the NATTS MDL MQO or 10<sup>-6</sup> cancer risk except acrolein (3 ppt) and 1,1,2,2-tetrachloroethane (~10-fold).
- This indicates laboratories are sufficiently sensitive to meet NATTS MDL MQOs or cancer risk metrics.

VC	DCs (	ppbv)
	typical MDL	MDL MQO or 10 <sup>-6</sup> cancer risk
Acrolein	0.042	<u>0.039</u>
Benzene	0.024	<u>0.041</u>
1,3-Butadiene	0.024	<u>0.050</u>
Carbon tetrachloride	0.020	<u>0.027</u>
Chloroform	0.020	<u>0.100</u>
1,2-Dibromoethane	0.025	NA
1,2-Dichloroethane	0.026	NA
Dichloromethane	0.027	0.600
1,2-Dichloropropane	0.026	NA
cis- 1,3-Dichloropropene	0.020	0.066
trans-1,3-Dichloropropene	0.021	0.066
1,1,2,2-Tetrachloroethane	0.027	0.0025
<b>Tetrachloroethylene</b>	0.020	<u>0.025</u>
<b>Trichloroethylene</b>	0.020	<u>0.037</u>
Vinyl chloride	0.028	<u>0.043</u>



Metals (ng/m <sup>3</sup> )											
	Typica	I MDL									
	High Volume Metals	MDL MQO or cancer risk 10 <sup>-6</sup>									
Antimony	0.0094	0.028	20								
<u>Arsenic</u>	0.0231	0.106	<u>0.23</u>								
<u>Beryllium</u>	0.0014	0.034	<u>0.42</u>								
<u>Cadmium</u>	0.0020	0.014	<u>0.56</u>								
Cobalt	0.0048	0.013	10								
Lead	0.0069	0.067	<u>15</u>								
<u>Manganese</u>	0.0051	0.166	<u>5</u>								
<u>Nickel</u>	0.0051	0.293	<u>2.1</u>								
Selenium	0.0123	0.182	<u>2000</u>								

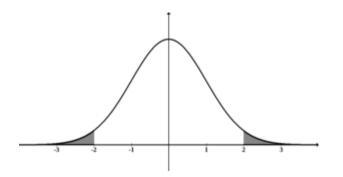
(ppbv)									
	<u>MDL</u> <u>MQO</u> or								
typical MDL	cancer risk 10 <sup>-6</sup>								
0.0254	<u>0.250</u>								
0.0253	NA								
0.0265	<u>0.064</u>								
0.0163	NA								
	typical MDL 0.0254 0.0253 0.0265								

Carbonyla

PAHs (ng/m <sup>3</sup> )										
	typical MDL	MDL MQO or cancer risk 10 <sup>-6</sup>								
Acenaphthene	0.222	300								
Anthracene	0.219	300								
<u>Benzo(a)pyrene</u>	0.347	<u>0.91</u>								
Fluoranthene	0.257	300								
Fluorene	0.237	300								
<u>Naphthalene</u>	0.473	<u>29</u>								
Phenanthrene	0.249	300								
Pyrene	0.228	300								



- NATTS data were pulled from AQS covering 2003-2010 and binned according to percentiles:
  - 5, 10, 25, 50, 75, 90, & 95%
- These binned data were compared to the typical MDLs to determine whether the specific percentile was above or below the typical MDL.





VOCs (ppbv)											
analyte	typical MDL	reported data (percentile)									
		5	10	25	50	75	90	95			
Acrolein	0.042	no data in table									
Benzene	0.024	0.15	0.26	0.41	0.64	1.00	1.59	2.09			
1,3-Butadiene	0.024	0.00	0.00	0.00	0.04	0.09	0.18	0.27			
Carbon tetrachloride	0.020	0.00	0.25	0.48	0.57	0.65	0.75	0.82			
Chloroform	0.020	0.00	0.00	0.00	0.10	0.18	0.34	0.59			
Tetrachloroethylene	0.020	0.00	0.00	0.00	0.09	0.20	0.40	0.61			
Trichloroethylene	0.020	0.00 0.00 0.00 0.04 0.08 0.11									
Vinyl chloride	0.028	0.00	0.00	0.00	0.00	0.00	0.02	0.03			

Green cells indicate concentrations less than the typical MDL.



5

0.000

7.806

0.347

0.473

Benzo(a)pyrene

Naphthalene

carbonyls (ppbv)												
analyte	typical MDL			reporte	d data (pe	rcentile)						
		5	10	25	50	75	90	95				
Acetaldehyde	0.03	0.44	0.57	0.85	1.30	2.00	2.99	3.77				
Formaldehyde	0.03	0.61	0.86	1.41	2.22	3.50	5.08	6.50				
	PAHs (ng/m <sup>3</sup> )											
analyte	typical MDL			reporte	ed data (po	ercentile)						

Green cells indicate concentrations less than the comparison MDL.

10

0.000

12.498

25

0.000

27.385

50

0.031

75

0.100

90

0.244

55.675 100.622 164.229 213.253

95

0.404



#### metals - high volume (ng/m<sup>3</sup>)

			U				<u> </u>	/
analyte	typical	ile)						
analyte	MDL	5	10	25	50	75	90	95
Arsenic	0.023	0.080	0.160	0.310	0.550	0.920	1.600	2.210
Beryllium	0.001	0.000	0.000	0.000	0.000	0.010	0.080	0.100
Cadmium	0.002	0.000	0.000	0.050	0.090	0.170	0.340	0.600
Lead	0.007	0.650	0.900	1.400	2.410	4.020	7.220	10.000
Manganese	0.005	0.920	1.240	2.020	3.640	7.900	18.400	28.390
Nickel	0.005	0.050	0.210	0.610	1.000	1.900	3.360	5.050

#### metals - low volume (ng/m<sup>3</sup>)

analyte	typical		re	ported	data (pe	rcentile	)	
analyte	MDL	5	10	25	50	75	90	95
Arsenic	0.106	0.000	0.060	0.240	0.450	0.770	1.160	1.480
Beryllium	0.034	0.000	0.000	0.000	0.000	0.010	0.030	0.040
Cadmium	0.014	0.000	0.000	0.040	0.070	0.120	0.220	0.340
Lead	0.067	0.520	0.800	1.370	2.330	4.130	6.950	9.420
Manganese	0.166	0.600	1.030	2.390	5.210	9.610	15.410	20.270
Nickel	0.293	0.000	0.000	0.340	0.860	1.740	5.200	10.300

Green cells indicate concentrations less than the comparison MDL.



- Blank data (lot blanks, field blanks, trip blanks, and lab blanks) were polled from AQS from 2003-2010 to determine a mean or median blank value for each analyte.
- To approximate the MDL<sub>b</sub> for each analyte in a worst case scenario, the typical MDL in the previous step was added to the mean or median blank value.







- In the following slides:
  - Green = less than the typical MDL and MDL<sub>b</sub>
  - Red = less than the MDL<sub>b</sub>
  - Orange = MDL<sub>b</sub> > MDL MQO



#### VOCs (ppbv)

analyte	typical MDI	blank value	MDL <sub>b</sub>	reported data (percentile)							
analyte											
				5	10	25	50	75	90	95	
Acrolein	0.042	not in AQS	0.042	no data in table							
Benzene	0.024	0.050	0.074	0.15	0.26	0.41	0.64	1.00	1.59	2.09	
1,3-Butadiene	0.024	0.000	0.024	0.00	0.00	0.00	0.04	0.09	0.18	0.27	
Carbon tetrachloride	0.020	0.003	0.023	0.00	0.25	0.48	0.57	0.65	0.75	0.82	
Chloroform	0.020	0.002	0.022	0.00	0.00	0.00	0.10	0.18	0.34	0.59	
Tetrachloroethylene	0.020	0.000	0.020	0.00	0.00	0.00	0.09	0.20	0.40	0.61	
Trichloroethylene	0.020	0.000	0.020	0.00	0.00	0.00	0.00	0.04	0.08	0.11	
Vinyl chloride	0.028	0.000	0.028	0.00	0.00	0.00	0.00	0.00	0.02	0.03	

Green cells indicate concentrations less than the comparison MDL.



carbonyls (ppbv)										
analyte	typical MDL	blank value	MDL <sub>b</sub>	reported data (percentile)						
				5	10	25	50	75	90	95
Acetaldehyde	0.025	0.038	0.063	0.44	0.57	0.85	1.30	2.00	2.99	3.77
Formaldehyde	0.026	0.052	0.079	0.61	0.86	1.41	2.22	3.50	5.08	6.50

PAHs (ng/m <sup>3</sup> )										
analyte	typical MDL	blank value	MDL <sub>b</sub>							
				5	10	25	50	75	90	95
Benzo(a)pyrene	0.347	0.008	0.355	0.000	0.000	0.000	0.031	0.100	0.244	0.404
Naphthalene	0.473	1.229	1.703	7.806	12.498	27.385	55.675	100.622	164.229	213.253

Orange cells indicate MDLs which exceed the NATTS MDL MQO. Green cells indicate concentrations less than the comparison MDL.



#### metals - high volume (ng/m<sup>3</sup>)

	analyte typical blank MDL value	blank		reported data (percentile)							
analyte			MDL <sub>b</sub>	5	10	25	50	75	90	95	
Arsenic	0.023	0.036	0.059	0.080	0.160	0.310	0.550	0.920	1.600	2.210	
Beryllium	0.001	0.003	0.004	0.000	0.000	0.000	0.000	0.010	0.080	0.100	
Cadmium	0.002	0.053	0.055	0.000	0.000	0.050	0.090	0.170	0.340	0.600	
Lead	0.007	0.302	0.309	0.650	0.900	1.400	2.410	4.020	7.220	10.000	
Manganese	0.005	0.757	0.763	0.920	1.240	2.020	3.640	7.900	18.400	28.390	
Nickel	0.005	0.446	0.451	0.050	0.210	0.610	1.000	1.900	3.360	5.050	

#### metals - low volume (ng/m<sup>3</sup>)

analyte	typical MDL	blank value	$MDL_b$	reported data (percentile)						
				5	10	25	50	75	90	95
Arsenic	0.106	0.090	0.196	0.000	0.060	0.240	0.450	0.770	1.160	1.480
Beryllium	0.034	0.004	0.037	0.000	0.000	0.000	0.000	0.010	0.030	0.040
Cadmium	0.014	0.035	0.048	0.000	0.000	0.040	0.070	0.120	0.220	0.340
Lead	0.067	0.046	0.113	0.520	0.800	1.370	2.330	4.130	6.950	9.420
Manganese	0.166	0.146	0.312	0.600	1.030	2.390	5.210	9.610	15.410	20.270
Nickel	0.293	0.259	0.552	0.000	0.000	0.340	0.860	1.740	5.200	10.300

Green cells indicate values less than the typical MDL and  $MDL_b$ . Red cells indicate values less than the  $MDL_b$ .

#### **Example MDL Scenario**

Lab Q determined their formaldehyde MDL by preparing and analyzing 8 spiked cartridges (spiked at 0.030 µg/cartridge) and 8 blanks.

aliquot	measured concentration (µg/cartridge)				
aliquot	Spikes	Blanks			
1	0.1685	0.1412			
2	0.1651	0.1399			
3	0.1701	0.1402			
4	0.1673	0.1405			
5	0.1692	0.1408			
6	0.1686	0.1403			
7	0.1705	0.1402			
8	0.1696	0.141			
average	0.1686	0.1405			
stdev	0.0017	0.0004			
student's T	2.998	2.998			
calculated MDL	0.0052	0.0013			
calculated MDL + blank average	NA	<u>0.1419</u>			
spike value/calculated MDL	5.8	NA			

The MDL<sub>b</sub> of 0.1419 is higher than the MDL<sub>sp</sub> of 0.0052  $\mu$ g/cartridge and is reported as the laboratory MDL (0.0985  $\mu$ g/m<sup>3</sup>).



# Summary

- With rare exception existing MDLs meet MDL MQOs.
- Draft NATTS TAD Revision 3 adopts the MUR.
- The MUR includes temporal variability and matrix background in the MDL determination.
- Under a worst case scenario, MDLs by the proposed MUR meet the MDL MQOs with the exception of formaldehyde, and with the exception of cadmium and nickel, no additional data would be flagged as < MDL for these methods.







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