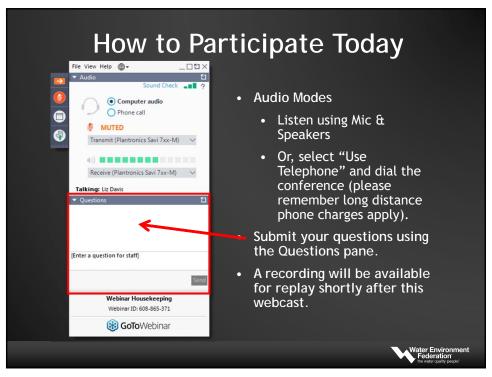


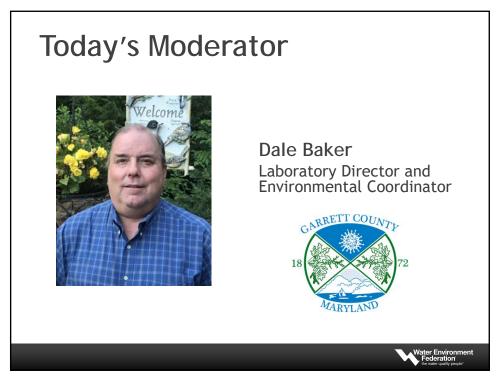
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# Calculating & Using Method Detection Limits

A Joint Presentation from Water Environment Federation & American Public Health Association







## Today's Speakers

- History of Method Detection Limits & Regulatory Programs
  - Elizabeth Turner
- Evaluating Blanks
  - Jeff Bennett
- Calculating MDLs
  - Mary Johnson



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## Our Next Speaker



Elizabeth Turner Quality Program Manager





## History of Method Detection Limits & Regulatory Programs

Elizabeth Turner



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## Why Method Detection Limits?

- Make quantitation meaningful
- Needed for risk assessment
  - Regulatory Programs
  - Statistical analysis

Protection of human health and the environment to a large degree depends on the ability to measure accurately the presence or absence of contaminants of concern.



### **Detection Limit**

Carson se tre low (uni

- Can you see me now? (12 pt)
- Can you see me now (18 pt)
- · Can you see me now? (24)

At what font size can you see words?



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### Analytical detection limits are:

- developed by statisticians,
- applied by analytical laboratories,
- and used by policy makers, regulators, and lawyers.
  - Michael Brisson



## Lloyd Currie - 1968

- Introduced terms of "critical level" (LC), "critical value" (CRV); the "detection decision"; with a 50% confidence level
- "minimum detectable value" (MDV), "detection limit" (LD) with a 99% confidence level
- "determination limit", "minimum quantifiable value" (MQV); limit of quantitation" (LOQ); commonly "quantitation limit" (LQ) required precision, accuracy, false negative error rate and qualitative identification criteria for the intended purpose.



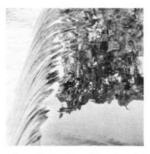
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# Trace analyses for wastewaters

Method detection limit, a new performance criterion for chemical analysis, is defined as that concentration of the analyte that can be detected at a specific confidence level. Both theory and applications are discussed for reliable wastewater analyses of priority pollutants

John A. Glaser
Denis L. Foerst
Gerald D. McKee
Stephan A. Quave
William L. Budde
U.S. Environmental Protection
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The development of trace analysis methodology brought with it a series of questions about method performance at low concentration levels of analyte (1, 2, 3). Under Section 304(h) of the Clean Water Act, as amended in 1977, (4) the Environmental Monitoring and



ority pollutants, it was incumbent on EMSL to develop method perfordetection limit should be related to the standard deviation of the measured values at or near zero concentration of the analyte (11).

There is no doubt that the detection limit is one of the most important performance characteristics of an analytical procedure. In most cases, a detection limit must be viewed as a temporary limit to current methodology.

#### Complete analytical system

Ostensibly, analysts do not directly observe concentrations of analyte. The measurements of the transducer signal, which are related to the analyte concentration, are actually observed. In any analytical system, information



### EPA - 1981

- Method detection limit (MDL) was first published in a paper by John Glaser and others at EPA's laboratory in Cincinnati in 1981 in Environmental Science and Technology
- MDL based on Currie's work
- Employs low-level spikes rather than backgrounds
- Uses Student's t-test to allow for varying number of replicates



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### **Various Procedures**

- American Council of Independent Laboratories (ACIL) Proposed Procedures for Determining the Method Detection Limit and Quantitation Limit (ACIL procedure)
- Hubaux-Vos Detection Limit Procedure
- ASTM Interlaboratory Detection Estimate (IDE)
- EPA MDL, 40 CFR Part 136, Appendix B
- ASTM Interlaboratory Quantitation Estimate (IQE)
- EPA OGWDW Lowest Concentration Minimum Reporting Level (LC-MRL) for Quantitation
- ISO/IUPAC
- USGS LT-MDL



### **EPA Method Detection Limit**

- 1984 2017
- 40 CFR 136 Appendix B
- The method detection limit (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.



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### **Procedure**

- 7 low level spikes
- 2 10x the expected MDL
- $MDL = T_{(n-1,1-\alpha=0.99)}(S)$
- Performed annually
- Usually done under ideal conditions



### **Limitations of MDL Procedure 1.11**

- Assumption of normal distribution and constant standard deviation
- Narrow estimate of method variability
- Assumption instrument variability is constant
- Assumption variability is the same for all instruments
- · Assumption of normal distribution
- Focused on false negatives, ignored false positives



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## **EPA Must Change Procedure**

- In 1999, several industry groups filed suit against EPA (Alliance of Automobile Manufacturers, et al. v. EPA, No. 99-1420, (D.C. Cir.)) - re: EPA Method 1631E
- October 2000, the parties reached a settlement agreement that required EPA to assess existing Agency and alternative procedures for determining detection and quantitation limits and sign a notice for publication in the Federal Register on or before February 28, 2003, and to invite comment on the assessment.



## **EPA Must Change Procedure**

- 2002 USEPA issues a Technical Support Document of Detection and Quantitation Regulations under the Clean Water Act (TSD).
- 2003 Draft revised MDL published
- 2003 Consensus letter submitted to Assistant Administrator of Office of Water signed by 31 parties urging EPA to consider a scientifically sound approach to the detection and quantification issue.
- 2004- proposed MDL procedure was withdrawn



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## Federal Advisory Committee

- 2004 Federal Register notice published that a neutral party is seeking a broad group of stakeholders willing to work together to define and address concerns about the way detection and quantitation values are calculated and used to support CWA programs.
- Formed in 2005
- Composed of state government, environmental laboratories, regulated industry, public utilities, the environmental community, and EPA
- To provide advice and recommendations on approaches for the development of detection and quantitation procedures and uses of these procedures in Clean Water Act program



### **Committee Recommendations**

December 2007 - 196 page Committee report

- A ≤ 1% false positive rate be used for detection.
- Need for Detection Limit and Quantitation Limit estimates that reflect normal, routine operations.
- Ongoing verification of detection limit and quantitation limit



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## **EPA Must Change Procedure**

- 2009 EPA Pilot study for 200.7 and 625 concluded that additional data generated using other analytical methods and more laboratories are needed to fully assess the applicability of these procedures to Clean Water Act Programs
- 2010 TNI forms Environmental Methods Measurement Expert Committee based on a USEPA grant to address Calibration, Detection, Quantification and other measurement issues.
- 2013 TNI EMEC (renamed Chemistry committee) completes work on a MDL revision and submits to EPA
- 2014 EPA completes internal review of the revised MDL and makes minor modifications
- 2015 EPA publishes revised MDL as part of a Methods Update Rule
- 2017 Signed by EPA Administrator Scott Pruitt



### **EPA MDL Procedure 2**

- 2017 Method Update Rule
- 40 CFR 136 Appendix B
- The method detection limit (MDL) is defined as the minimum measured concentration of a substance that can be reported with 99% confidence that the measured concentration is distinguishable from method blank results.



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## MDL Changes

- Initial study over 3 days
- Use of blanks
- The MDL now requires that the samples used to calculate the MDL are representative of laboratory performance throughout the year, rather than on a single date (MDLv)
- A laboratory has the option to pool data from multiple instruments to calculate one MDL that represents multiple instruments. (Not for Drinking Water)
- Recalculate every 13 months



## MDL Influenced By

- Sample Matrix
- Preparation Steps
- Instrument (age, maintenance)
- Technology (GC-MS ≠ GC-FID)
- Analyst Skill
- Environmental Conditions



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### **EPA MDL Procedure 2**

- Analyze 7 blanks and 7 blanks prepared and analyzed in at least 3 batches over 3 separate days.
- Multiple Instruments
  - A minimum of two spiked samples and two method blank samples prepared and analyzed on different calendar dates.
- Calculate MDL<sub>s</sub> and MDL<sub>b</sub>
- MDL is higher of MDL<sub>s</sub> and MDL<sub>b</sub>
- TNI Tip use LOQ for MDL spike



## **MDL Verification Samples**

- During any quarter in which samples are being analyzed, prepare and analyze a minimum of 2 spiked samples on each instrument, in separate batches, using the same spiking concentration used for the initial MDL study.
- Evaluate MDL, against acceptance criteria
- Ensure that at least 7 spiked samples and 7 method blanks are completed for the annual verification.
- Missed a verification?

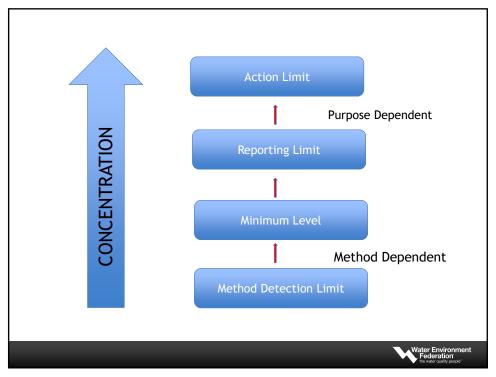


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### **Annual Verification**

- Every 13 months recalculate MDL<sub>s</sub> and MDL<sub>b</sub>
  - · Data from last 24 months
  - MDL verification spikes and method blanks
- The verified MDL is the greater of the  $MDL_s$  or  $MDL_b$  .
- If the verified MDL is within 0.5 to 2.0 times the existing MDL, and fewer than 3% of the method blank results have numerical results above the existing MDL, then the existing MDL may optionally be left unchanged





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### Minimum Level

- The lowest level at which the entire analytical system must give a recognizable signal and acceptable calibration point for the analyte. It is equivalent to the concentration of the lowest calibration standard, assuming that all methodspecified sample weights, volumes, and cleanup procedures have been employed.
- The ML is calculated by multiplying the MDL by 3.18 and rounding the result to the number nearest to (1, 2, or 5) x 10n, where n is an integer.
- Minimum levels are used in some US EPA methods.



## **Reporting Limits**

- May or maynot be equal to quantitation limit
- Lowest standard must be at or below limit
- · No federal method for determination
- Many states have own process for establishing for regulatory programs



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## Regulatory Reporting Limits

Allowable concentration levels for a contaminant in a particular environmental medium (e.g., soil, air, or water) are often based on health-related risk assessments and are sometimes lower than levels that can be quantitated in a laboratory. For this reason, the lowest quantifiable level frequently becomes the de facto regulatory limit for monitoring and compliance purposes.



### **Use in NPDES Permits**

- MDLs serve as base for Reporting Limit
- Protective of Water Quality Standards
- State defined reporting limits
- Must use sufficiently sensitive method



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## Reporting Limits for NPDES

- Southwest state 3x MDL of most sensitive method for analyte or minimum level
- Central State- 5X method MDL
- Eastern state 5X method MDL
- Eastern State report everything to MDL



## Other Regulatory Uses

- Ambient monitoring (305)
- Pretreatment and Stormwater
- · Water Quality Criteria
- 303(d) listing for Total Maximum Daily Loads
- Drinking Water monitoring (MCLs and reporting limits)
- Remediation (Protective Concentration Levels)



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## **Key Take Aways**

- Detection ≠ Quantitation
- Detection limits will vary by laboratory
- Detection limits are utilized by regulators to assess:
  - Establish Permit Limits
  - Risk
  - Compliance



## Our Next Speaker



Jack Bennett Technical Manager, Analytical Laboratory





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## **Evaluating Blanks**

Jack Bennett





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### **Gathering Data**

- The MDL must be calculated in the units that are reported for samples.
  - Not the units from the calibration curve.
  - Use the nominal sample weight or volume and take through all calculations.
- The results used for the MDL calculation must not be censored.
  - This really applies to blanks, although it could apply if the MDL spike is around the reporting limit.
  - Can't use "less than" or "Not Detected" or "zero" if the instrument gives a numerical result.
    - This is most common for metals, methods with "common" blank contaminants, and some automated wet chem methods.





### **Gathering Data**

- If you have a LIMS and normally do not report below the Reporting Limit (RL), create a LIMS test code without the logic to censor the data.
- If you don't have a LIMS, you can use a spreadsheet to do the calculations.
  - Many instruments have an option to export a file, which can be used to populate a calculations spreadsheet.
    - In Excel, the Legacy Wizard is (in my opinion) more user friendly than the new Wizard.
  - Its very important to keep up with "filing" the data as it is generated rather than gathering it once a year.
    - Especially important for blanks.

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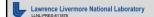
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	Dates Run	7/24/2018	7/24/2018	7/24/2018	7/26/2018	7/26/2018	7/26/2018	8/2/2018	8/2/2018								
	Spike											Spike/MDL		RL/MDL >2*		Initial MDL,	RL/MDL >3***
lement b 206.836	Level, ug/g	201812385A 4.9700	201812385B 4.9800	4.3000	5.7300	5.4300	201812385F 5.5400	201812385G 6.9700	201812385H 5.6000	0.7739	ug/g 2.320	<10 ** 2.15	RL,ug/g 7.5	3.23	150%* 108.80	ug/g 2.456	3.23
s 193.696	5		5.4000	4.8000	5.7100	5.5700	4.7500		4.7400	0.7739	1.572	3.18				1.648	4.77
a 233.527 Rad	4		3.4000	3.7300		4.1900	3,7000		3.7600	0.3244	0.761	5.26		5.26	94.75	0.971	5.26
e 313.107	0.04			0.0355		0.0381	0.0407		0.0419	0.0043	0.013	3.08			100.66	0.017	38.47
d 214.440	0.3	0.3110	0.3200	0.3160		0.3030	0.2900		0.3080	0.0120	0.036	8.35	0.3	8.35	102.96	0.036	8.35
r 205.560	2.5	2.4800	2.3900	2.4500	2.5100	2.4600	2.4600		2.5100	0.0391	0.117	21.34	2.5	21.34	98.75	0.117	21.34
o 228.616	2.4	2.4600	2.4000	2.4100	2.5000	2.4000	2.4200	2.5300	2.5000	0.0520	0.156	15.39	2.4	15.39	102.19	0.156	15.39
u 324.752	12	12.3000	12.4000	12.3000	12.8000	12.3000	12.2000	12.7000	12.7000	0.2326	0.697	17.21	18	25.81	103.85	0.697	25.81
b 220.353	1.6		1.8400	1.7000		1.6000	1.6600		1.7000	0.1347	0.404	3.96		4.46	102.73	0.404	4.46
to 202.031	4		3.9300	3.9900	4.1100	3.9000	3.8900		3.9200	0.0725	0.217	18.40	4	18.40	99.09	0.217	18.40
1 231.604	10		10.6000	10.6000		10.5000	10.6000		10.7000	0.1195	0.358	27.91	10	27.91	106.50	0.358	27.91
e 196.026	- 6		5.5000	7.1800		5.5000	4.5000		6.0900	1.0417	3.123	1.92		6.40	89.02	3.123	6.40
g 328.068	1.92		0.5000	0.4930	0.5660	0.5760	0.5900		0.6880	0.0770	0.231	8.32		8.32	30.42	0.231	8.32
1 190.801	1 10		0.5650 9,4400	1.0000 9.4000	0.8100	1.0600 9.6400	0.4920 9.5800		1.0800	0.2453 0.3471	0.735 1.041	1.36 9.61	7 10	9.52	89.59 96.84	0.919	9.52
n 213.857	10		11,9000	11.7000		9.6400	9.5800	11.1000	11,2000	0.34/1	0.945	10.59	10	10.59	113.75	1.041	9.61
11 213.037	10	11.0000	11.9000	11.7000	11.3000	11.1000	11.1000	11.1000	11.2000	0.3131	0.943	10.59	10	10.59	115.75	1.837	10.59
=Not required to																	
*= Use as guidano		ining ruture N	TUL SPIKING Te	veis													
**NELAC 2016 Re	quirement																

### The Blank MDL – Why?

- The original MDL method was based on an assumption that the blanks were essentially zero.
  - Normal distribution tightly around zero.
- Without going into too much detail, Lloyd Currie's paper "Limits for Qualitative Detection and Quantitation: Application to Radiochemistry" Anal. Chem. 40, 586-593 (1968) was the seminal paper on the concept of detection limits.
- He proposed that the point where there was a random chance of a false positive being ≤ 1% was the Critical Level (L<sub>c</sub>).





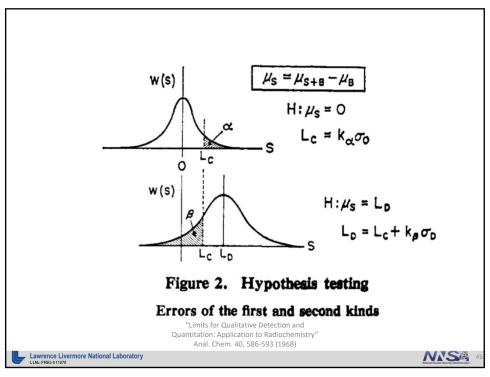
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### The Blank MDL – Why?

- The concentration that gave that point was the Detection Limit  $(L_D)$ .
- This figure from his paper illustrates the difference between the blank population and the detection limit population:

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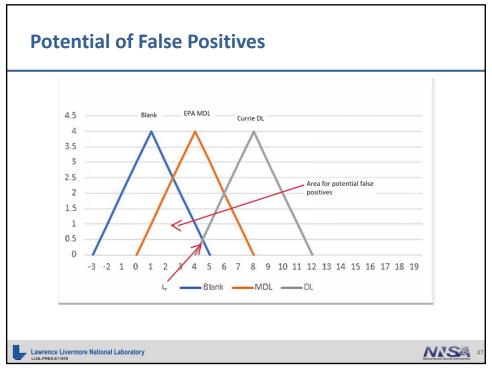
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### The Blank MDL – Why?

- Curries approach was to minimize false positives.
- When EPA produced their original MDL procedure in 1984, their approach was to minimize false negatives.
  - They set their procedure so that the MDL was at the L<sub>c</sub>.
    - Blank < MDL/L<sub>c</sub> < Currie L<sub>D</sub>
- Blanks were not included in the EPA MDL calculation.
- As methods got more sensitive, labs were reporting false positives.
  - False positives can have consequences.

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### The Blank MDL – Why?

- In 1999, EPA was sued over a low-level mercury MDL method and lost.
- It took until 2016 for them to figure out a new MDL procedure.
  - Why not just use Curries procedure?
    - It is not practicable for a typical lab.
      - Need to run lots of samples.
- Including blanks in the procedure is a way to reduce false positives when reporting results below the RL.
- Not perfect, but not going to change.
- What else can use for solids and other methods that can't practically be spiked.
  - Not a requirement, but (maybe) a "best practice".



NNSA 4

#### Blanks in the MDL Rev 2

- Initial MDL for Method Blanks:
  - Can use routine Method Blanks.
    - Must be within the last 24 months.
  - If no routine Method Blanks, at least 7 Method Blanks processed through the entire sample prep and analysis process on three separate calendar days.
    - If multiple instruments are used, must be run on all instruments.
    - A minimum of two blanks prepared on different days is required for each instrument.
    - Statistical outlier removal procedures should not be used.

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### Calculating the Initial Blank MDL (MDL<sub>h</sub>)

- Three options:
  - Option 1 No method blanks give numerical results, the MDL<sub>b</sub> does not apply.
    - A negative number as a result is a numerical result.
    - Results below the current MDL or RL are numerical results.
    - An example of a non-numerical result is a chromatography method when a peak is not present.
  - Option 2 Some (but not all) results give a numerical result, set the MDL<sub>b</sub> to the highest method blank result.
    - If using routine method blank data and there are >100 method blanks, set the  $MDL_h$  to  $99^{th}$  percentile.
      - Estimating the 99th percentile is acceptable.

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### Calculating the Initial Blank MDL (MDL<sub>b</sub>)

 Option 3 – All the method blanks give either positive or negative numerical results, calculate the MDL<sub>h</sub> as:

$$MDL_b = \bar{x} + t_{students} (S_b)$$

where:

 $MDL_b$  = the MDL based on method blanks.  $\bar{x}$  = mean of the method blank results.

(use zero for the mean if the mean is negative)

 $t_{students}$  = Students t-value for the 99<sup>th</sup> percentile. S<sub>b</sub> = Sample standard deviation.

 If existing data is being used and there are more than 100 method blanks, the 99th percentile value of the results can be used.

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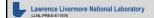
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				MDL Study EF	A M DL Proce	dure Revision	2						
				,	SDG								
tudents T Value	2.998			Instrument	ICP 6		Matrix	Soil/Solid		Method	EPA 3050B/6010B		
	Dates Run	7/24/2018	7/24/2018	7/24/2018	7/26/2018	7/26/2018	7/26/2018	8/2/2018	8/2/2018				4
	Spike										Blank MDL,		Average LRB,
lement	Level, ug/g	MDI BIK1	MDL BLK2	MDL BLK3	MDL BLK4	MDL BLK5	MDL BLK6	MDL BLK7	MDL BLK8	Std. Dev	ug/g	RL, ug/g	ug/g
b 206.836	0			-0.0450	0.8210	0.7700			1.1500			7.5	0.22
s 193.696	0			0.4180	0.1360	0.6470			0.8100	0.4483		7.5	0.30
a 233.527 Rad	0			-0.6210	0.0194	-0.1380						4	-0.25
e 313.107	0		-0.0059	-0.0008	-0.0100	-0.0014			-0.0014			0.5	0.00
d 214.440	0		-0.0100	-0.0136	-0.0080	-0.0239			-0.0037			0.3	-0.01
r 205.560	0			-0.0340	-0.0620	-0.0550			-0.0320			2.5	-0.04
o 228.616	0	-0.1400	-0.0479	-0.0391	0.0372	-0.0607	-0.0434	-0.0103	-0.0215	0.0503	0.151	2.4	-0.04
u 324.752	0	-0.0350	-0.1120	-0.1030	-0.1000	-0.1490	-0.1440	0.3630	0.1550	0.1817	0.545	18	-0.02
b 220.353	0		0.0930	0.1150	0.0010	0.1010	0.0410	0.0490	-0.0140	0.0691		1.8	0.07
1o 202.031	0			-0.0150	0.0190	-0.0470				0.0300		4	-0.02
i 231.604	0			-0.0010	0.0180	0.0020						10	0.02
e 196.026	0			-0.5620	0.6040	0.1860			-2.1500	1.0234		20	-0.32
g 328.068	0			-0.0380	-0.0300	-0.0560			0.0430			1.92	-0.03
190.801	0			-0.6820	-0.0410	-0.3700				0.3064		7	-0.35
292.402 in 213.857	0		-0.1520	-0.0446	0.1720	-0.1290						10	-0.08
	0	0.4500	1.5400	0.4800	0.4170	0.2030	0.2210	0.2120	0.0786	0.4626	1.837	10	0.45

### **Ongoing Verification for Method Blanks**

- The data only has to be collected if samples are being analyzed in a quarter.
  - There is guidance in the EPA MDL FAQ about low volume tests at <a href="https://www.epa.gov/cwa-methods/method-detection-limit-frequent-questions">https://www.epa.gov/cwa-methods/method-detection-limit-frequent-questions</a>
- The verification / recalculation is done every 13 months ideally using data from the last 24 months.
  - Only use data from acceptable batches.
  - The 99<sup>th</sup> percentile value is not listed as an option for ongoing verification of methods with lots of blanks, however:
  - "The laboratory has the option to use only the last six months of method blank data or the fifty most recent method blanks, whichever criteria yields the greater number of method blanks".
- Must use data from <u>all</u> acceptable batches.





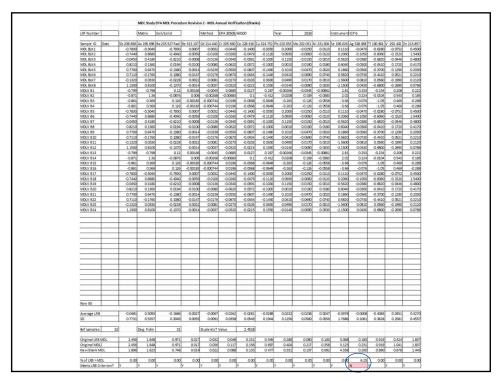
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### **Ongoing MDL Verification Criteria**

- Fewer than 3% of the method blank results can have results greater than the existing MDL.
  - If more than 3% of the method blanks are greater than the existing MDL,
     must use the new verification MDL.
- MDL Evaluation Criteria:
  - If the verified MDL (i.e. MDL calculated using the verification data) is:
    - Within 0.5 to 2 times the existing MDL.
    - Fewer that 3% of the method blank results are above the existing MDL.
  - Then the existing MDL may continue to be used.
  - Otherwise, use the newly calculated verification MDL.
    - $\bullet\,$  The  $\,$  verification MDL is the greater of the method blank MDL or the Spike MDL
  - However, if more than 5% of the MDL Verification Spikes do not return positive numerical results, then the original MDL must be re-performed using a higher spiking level.

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			MDL Study	EPA MDL Proce	dure Revisio	n 2 - MDL A	nnual Verif	ication (Spil	es)								
LRF Number			Matrix	Soil/Solid		Method	EPA 3050B	/6010D		Year	2019		Instrumen	ICP 6			
Sample ID	Date	Sb 206.836			Be 313.107		Cr 205.560				Mo 202.031						Zn 213.8
MDL 1		4.970	4.400	4.000	0.044	0.311	2.480	2.460	12.300	1.690	3.960	10.600	5.640	0.557	1.110	9.210	11.6
MDL 2		4.980	5.400	3.320	0.038	0.320	2.390	2.400	12.400	1.840	3.930	10.600	5.500	0.500	0.565	9.440	11.9
MDL 3		4.300	4.800	3.730	0.036	0.316	2.450	2.410	12.300	1.700	3.990	10.600	7.180	0.493	1.000	9.400	11.
MDL 4		5.730	5.710	3.730	0.036	0.297	2.510	2.500	12.800	1.590	4.110	10.900	4.250	0.566	0.810	10.200	11.3
MDL 5		5.430	5.570	4.190	0.038	0.303	2.460	2.400	12.300		3.900	10.500	5.500	0.576	1.060	9.640	11.1
MDL 6		5.540	4.750	3.700	0.041	0.290	2.460	2.420	12.200	1.660	3.890	10.600	4.500	0.590	0.492	9.580	11.
MDL 7		6.970	5.750	3.890	0.048	0.326	2.490	2.530	12.700	1.370	4.010	10.700	4.070	0.703	1.050	10.000	11.1
MDL 8		5.600	4.740	3.760	0.042	0.308	2.510	2.500	12.700	1.700	3.920	10.700	6.090	0.688	1.080	10.000	11.2
MDLV 1		4.600	6.370	4.170	0.054	0.308	2.470	2.550	12.300	1.910	3.960	10.700	5.120	2.090	1.220	10.100	11.4
MDLV 2		4.470	4.340	4.350	0.041	0.338	2.680	2.620	13.000		4.120	10.800	5.320	2.090	1.550	10.300	11.8
MDLV 3		4.600	6.370	4.170	0.054	0.308	2.470	2.550	12.300	1.910	3.960	10.700	5.120	2.090	1.220	10.100	11.4
MDLV 4		4.470	4.340	4.350	0.041	0.338	2.680	2.620	13.000	1.740	4.120	10.800	5.320	2.090	1.550	10.300	11.3
MDLV 5		5.25	4.98	4.24	0.0401	0.318	2.44	2.47	12.9	1.64	3.93	10.7	3.01	2.08	1.93	10.3	9
MDLV 6		4.970	4.400	4.000	0.044	0.311	2.480	2.460	12.300	1.690	3.960	10.600	5.640	0.557	1.110	9.210	11.0
MDLV 7		4.980	5.400	3.320	0.038	0.320	2.390	2.400	12.400	1.840	3.930	10.600	5.500	0.500	0.565	9.440	11.9
MDLV 8		4.300	4.800	3.730	0.036	0.316	2.450	2.410	12.300	1.700	3.990	10.600	7.180	0.493	1.000	9.400	11.
MDLV 9		5.730	5.710	3.730	0.036	0.297	2.510	2.500	12.800	1.590	4.110	10.900	4.250	0.566	0.810	10.200	11.3
MDLV 10		5.430	5.570	4.190	0.038	0.303	2.460	2.400	12.300	1.600	3.900	10.500	5.500	0.576	1.060	9.640	11.1
MDLV 11		5.540	4.750	3.700	0.041	0.290	2.460	2.420	12.200	1.660	3.890	10.600	4.500	0.590	0.492	9.580	11.1
MDLV 12		6,970	5,750	3.890	0.048	0.326	2.490	2.530	12,700	1.370	4.010	10.700	4.070	0.703	1.050	10.000	11.1
MDLV 13		5,600	4,740	3,760	0.042	0.308	2.510	2.500	12,700	1.700	3.920	10.700	6.090	0.688	1.080	10.000	11.2
MDLV 14		4,600	6.370	4,170	0.054	0.308	2,470	2.550	12.300	1.910	3.960	10.700	5.120	2.090	1.220	10.100	11.4
MDLV 15		4 470	4 340	4,350	0.041	0.338	2,680	2.620	13.000	1.740	4,120	10.800	5.320	2.090	1.550	10.300	11.3
MDLV 16		4.9700	4.4000	4.0000	0.0442	0.3110	2.4800	2.4600	12.3000		3.9600	10.6000	5.6400	0.5570	1.1100	9.2100	11.6
Spike Level, ug/g		5	5		0.04	0.3	2.5	2.4	12	1.6	4	10	6	1 92	1	10	
spike Level, ug/g		,	,	,	0.04	0.3	2.3	2.4	12	1.0	- 4	10	,	1.92	- 4	10	
Std Dev		0.7234	0.6876	0.2962	0.0058	0.0136	0.0781	0.0731	0.2828	0.1388	0.0782	0.1073	0.9452	0.7019	0.3506	0.3908	0.50
# of samples	24		Deg. Frdm.	23		Students T	Value	2.4998									
0.1.1.14401		0.450	4.610			0.000	0.440	0.450	0.000			0.050	0.400			4.044	
Original MDL		2.456	1.648	0.971	0.017	0.036	0.117	0.156	0.697	0.404	0.217	0.358	3.123	0.231	0.919	1.041	1.
New Spike MDL Verification MDL		1.808 2.158	1.719 1.897	0.741	0.014	0.034	0.195	0.183	0.707	0.347	0.195	0.268	2.363 5.485	1.755	0.876 1.066	0.977	1.5
0o >95% of the					1	1			l				1	1	i l		
spikes return a					1	1			l				1	1	i l		
positive numerical																	
esult?		Υ	Υ	Y	Υ	Υ	Υ	Υ	Y	Υ	Υ	Y	Υ	$\sim$	Υ	Υ	Υ
s the original MDL /erified?		Y	Y	Y	Y	l <sub>Y</sub>	Y	Y	Y	Y	Υ	Y	l <sub>y</sub> /	N )	, I	Y	ΙΥ
Jse Verification																	

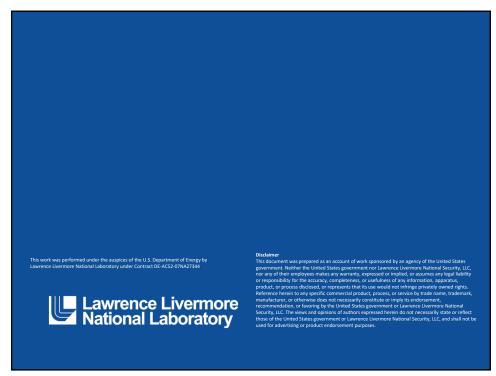
### **Takeaways:**

- Develop a system to keep track of when MDLV's are due.
  - This is especially important for low volume tests.
- Automate the calculations as much as possible.
  - Spreadsheets work nicely, and there is more than one way to get to your desired result.
    - Figure a way that makes sense to you.
  - Remember to (ideally) have someone else check your calculations.
- Start with an easy test.

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## Our Next Speaker



Mary Johnson Laboratory Manager



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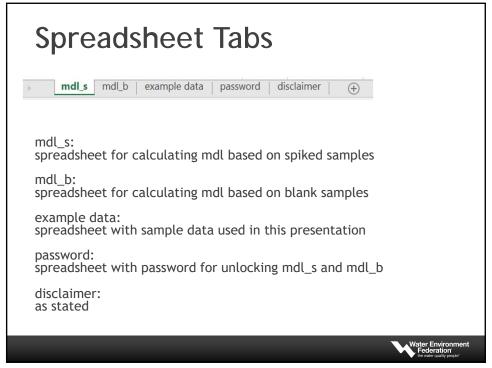
## Calculating MDLs

Mary Johnson



#### A Tool for Calculating MDLs Analyte: Analyte Name Spike Conc: (spike concentration must be a numerical value) Units: units Method: Method Reference or SOP Analysis Replicate Recovery units mg/L #DIV/0! #DIV/0! mg/L Average #DIV/0! #DIV/0! Std Dev #DIV/0! #DIV/0! Deg of Freedom MDL<sub>s</sub> = MDL based on spiked samples #NUM! MDL<sub>b</sub> = MDL based on blanks 0.016 MDL is greater of MDLs and MDLb #NUM!

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### A Disclaimer

These spreadsheets were put together by members of the Association of Public Health Laboratories and the WEF Laboratories Practices Committee. The authors have attempted to align procedures with the EPA's *Definition and Procedure for the Determination of the Method Detection Limit, Revision 2 (EPA821-R-16\_006)*.

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### **MDL Calculation**

The MDL is the greater of MDL, and MDL,

$$MDL_{s} = t_{(n-1, t-\alpha=0.99)} * S_{s}$$

$$MDL_{b} = X + (t_{(n-1, t-\alpha=0.99)} * S_{b})$$

You need a minimum of 7 spiked samples for the  $MDL_s$  calculation and 7 blank samples for the  $MDL_b$  calculation.



## $MDL_s$

$$MDL_s = t_{(n-1, t-\alpha=0.99)} * S_s$$

Where

MDLs = the method detection limit based on spiked sample

 $t_{(n-1, t-\alpha=0.99)}$  = Students t-value at 99% for standard deviation with

n - 1 degrees of freedom

S<sub>s</sub> = standard deviation of the spiked samples



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## Data Needed: Spiked Samples

- · Minimum of seven spiked samples
- · Must use most recent available data
- Data must be from at least three separate batches analyzed on three separate days
- Data must have been generated within last 24 months
- Analysis results must be a numerical value greater than zero
- No statistical outlier data removal for initial MDL



## Spiked Samples: Practicalities

- Spiking level is typically 2 10 times the expected MDL
- Analyzing two MDL samples each quarter is a practical way to collect enough data to calculate the MDL each year.
- Use only data associated with acceptable calibration and batch QC.



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## Example Data: MDL<sub>s</sub>

A laboratory tests for ammonia using a specific ion meter. Each quarter they analyze two 0.100 mg/L ammonia samples. The results of these analysis are used to calculate MDLs.

test date	mg/L Ammonia
1/1/2019	0.095
2/1/2019	0.091
4/1/2019	0.087
5/1/2019	0.088

test date	mg/L Ammonia
7/1/2019	0.104
8/1/2019	0.095
10/1/2019	0.088
11/1/2019	0.096



## It's Easier with a Spreadsheet

Analyte:	Ammonia			
Spike Conc:	0.100 mg/L			
Method:	SOP 301, Ammor	nia by Specific Ion	Electrode	(SM 4500-NH3 D)

	Test	Analysis		Percent
Replicate	Date	Result	units	Recovery
1			mg/L	0.0
2			mg/L	0.0
3			mg/L	0.0
4			mg/L	0.0
5			mg/L	0.0
6			mg/L	0.0
7			mg/L	0.0
8			mg/L	0.0
Average		#DIV/0!		0.0
Std Dev		#DIV/0!		0.0
Deg of Freedom		-1		
t(n-1)		#NUM!		
$MDL_s = MDL$ base	sed on spiked samp	oles		#NUM!

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## MDL<sub>s</sub> Calculation

#VALUE!	A
	Analysis
	Result
1	
2	
3	
4	
5	
6	
7	
8	
Average =	#DIV/0!
Std Dev =	#DIV/0!
Deg of Freedom =	-1
t(n-1) =	#NUM!
MDL =	#NUM!

Average: =average(A1:A8)

Standard Deviation = stdev(A1:A8)

(n - 1) Degrees of Freedom =count(A1:A8)-1

Students  $t_{(n-1, t-\alpha=0.99)}$ =ABS(TINV(2\*0.99,A9))

 $\frac{MDL}{MDL_s} = t_{(n-1, t-\alpha=0.99)} * S_s$ = A12 \* A10



## MDL<sub>s</sub> Calculation

	A
	Analysis
	Result
1	0.095
2	0.091
3	0.087
4	0.088
5	0.104
6	0.095
7	0.088
8	0.096
Average =	0.093
Std Dev =	0.006
Deg of Freedom =	7
t(n-1) =	2.998
$MDL_s =$	0.017

In this example, the MDLs calculation produced an MDLs of 0.017 mg/L.

$$MDL_s = t_{(n-1, t-\alpha=0.99)} * S_s$$
= 2.998 \* 0.006
= 0.017

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### Where to find Students t values

EPA's Method Detection Limit Publication: EPA 821-R-16-006

NIST.gov website:

https://www.itl.nist.gov/div898/handbook/eda/section3/eda3672.htm

Or just use the spreadsheet function for Students t.



## Organize Your Information

Analyte:	Ammonia			
Spike Conc:	0.100 mg/L			
Method:	SOP 301, Ammor	nia by Specific Ion	Electrode	(SM 4500-NH3 D)
	Test	Analysis		Percent
Replicate	Date	Result	units	Recovery
1	1-Jan-19	0.095	mg/L	95.0
2	1-Feb-19	0.091	mg/L	90.5
3	1-Apr-19	0.087	mg/L	87.0
4	1-May-19	0.088	mg/L	88.0
5	1-Jul-19	0.104	mg/L	104.0
6	1-Aug-19	0.095	mg/L	94.6
7	1-Oct-19	0.088	mg/L	88.0
8	1-Nov-19	0.096	mg/L	96.0
Average		0.093		92.9
Std Dev		0.006		5.7
Deg of Freedom		7		
t(n-1)		2.998		
$MDL_s = MDL$ ba	sed on spiked sam	ples		0.017

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### Is Your MDL Reasonable?

Is the calculated MDL > 0?

Is the calculated MDL > 0.1 \* spike concentration?

Is the spike concentration > calculated MDL?

Is the spike concentration between 1 and 10 times the MDL?



### Is Your MDL Reasonable?

In our example, for ammonia by specific ion electrode:

Spike Concentration = 0.100 mg/LAverage of eight replicates = 0.093Standard deviation = 0.006MDL<sub>s</sub> = 0.017



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### Is Your MDL Reasonable?

✓ Is the calculated MDL > 0?

0.017 > 0

✓ Is the calculated MDL > 0.1 \* spike concentration?

0.017 > 0.1 \* .100

✓ Is the spike concentration > calculated MDL?

0.100 > 0.017

✓ Is the spike concentration between 1 and 10 times the MDL?

0.017 < 0.100 < 0.17



# Do your spike results make sense?

	Percent
Replicate	Recovery
1	95.0
2	90.5
3	87.0
4	88.0
5	104.0
6	94.6
7	88.0
8	96.0
Average	92.9
Std Dev	5.7
Deg of Freedom	
t(n-1)	

Does the % recovery for each replicate make sense?

Are all spike results within <u>+</u> 2 standard deviations of the mean?



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## We're not done yet.

Remember: the MDL is the greater of  $\text{MDL}_{\text{s}}$  and  $\text{MDL}_{\text{b}}$ 

We still need to calculate MDL<sub>b</sub>



### Data Needed: Method Blanks

- Minimum of seven method blanks
- Must use most recent available data
- Data must be from at least three separate batches analyzed on three separate days
- Data must have been generated within the last 24 months.



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# How should we evaluate method blank data?

- If none of the method blanks give numerical results, MDL<sub>b</sub> does not apply
- If some, but not all, of method blanks give numerical results, MDL<sub>b</sub> is the highest method blank result. If more than 100 method blanks, set MDL<sub>b</sub> to number no less than 99<sup>th</sup> percentile.
- If all method blanks have numerical results,  $MDL_b = X + (t_{(n-1. t-\alpha=0.99)} *S_b)$



## $MDL_b$

$$MDL_b = X + (t_{(n-1, t-\alpha=0.99)} *S_b)$$

Where

MDL<sub>b</sub> = the method detection limit based on blank samples

X = mean of the method blank results

 $t_{(n-1, t-\alpha=0.99)}$  = Student's t-value at 99% for standard deviation with n-1

degrees of freedom

S<sub>b</sub> = standard deviation of the method blank analyses



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## Example Data: MDL<sub>b</sub>

Continuing our ammonia analysis example, let's assume the laboratory tested ammonia twelve times the previous year and thus have 12 blank results.

test date	mg/L Ammonia
1/1/2019	0.0029
2/1/2019	0.0123
3/1/2019	0.0000
4/1/2019	0.0060
5/1/2019	0.0071
6/1/2019	0.0071

test date	mg/L Ammonia	
7/1/2019	0.0069	
8/1/2019	0.0109	
9/1/2019	0.0058	
10/1/2019	0.0087	
11/1/2019	0.0023	
12/1/2019	0.0054	



## MDL<sub>b</sub> Calculation

Α	В	C	D
test	analysis		
date	result		
		average =	#DIV/0!
		S <sub>b</sub> , std dev =	#DIV/0!
		count =	0
		deg of freedom =	-1
		students $t_{(n-1)} =$	#NUM!
		MDL <sub>b</sub> =	$X + t_{(n-1)}(S_b)$
		=	#DIV/0!
	test	test analysis	test date         analysis result           average = Sb, std dev = count = deg of freedom = students t <sub>(n-1)</sub> =

Average:

=IF(average(B1:B12)<0, 0,average(B1:B12))

Standard Deviation
=stdev(B1:B12)

Count:
=count(B1:B12)

Degrees of Freedom =D3-1

Students t =ABS(TINV(2\*0.99,D4))

 $\begin{array}{l} \underline{MDL_b} \\ = X + (t_{(n-1, \ t-\alpha=0.99)} * S_b) \\ = D2 + (D3 * D4) \end{array}$ 



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## MDL<sub>b</sub> Calculation

test	analysis		
date	result		
1/1/2019	0.00290	average =	0.0062
2/1/2019	0.01230	S <sub>b</sub> , std dev =	0.003
3/1/2019	0.00000	count =	12
4/1/2019	0.00600	deg of freedom =	11
5/1/2019	0.00710	students $t_{(n-1)} =$	2.718
6/1/2019	0.00580		
7/1/2019	0.00690	MDL <sub>b</sub> =	$X + t_{(n-1)}(S_b)$
8/1/2019	0.01090	=	0.016
9/1/2019	0.00580		
10/1/2019	0.00870		
11/1/2019	0.00230		
12/1/2019	0.00540		

In this example, the  $\mathrm{MDL_b}$  calculation produced an  $\mathrm{MDL_b}$  of 0.016 mg/L.



### So what is the MDL?

The MDL is the greater of MDL<sub>s</sub> and MDL<sub>b</sub>

For our example:

 $\mathsf{MDL}_\mathsf{s} = 0.017$ 

 $MDL_b = 0.016$ 

So our MDL = 0.017 mg/L



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### MDL vs. RL

MDL - Method Detection Limit

RL - Reporting Limit

The RL is the smallest concentration that is reported by a laboratory. The RL may be the lowest standard used when making a calibration curve.



## References

Definition and Procedure for the Determination of the Method Detection Limit, Revision 2 EPA 821-R-16-006

Federal Advisory Committee on Detection and Quantitation Approaches and Uses in Clean Water Act Programs Final Report 12/28/07

ANALYTICAL DETECTION LIMIT GUIDANCE & Laboratory Guide for Determining Method Detection Limits Wisconsin DNR, PUBL-TS-056-96



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### Thank You!

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