

**Montana DEQ - Waste Management and Remediation Division**

**Data Validation Summary Form (Version 1.3.0, Revised 1/26/18)**

Please fill out the information below, using one form for each lab batch (one form can be used for multiple analytical methods). The form will grow and adjust, based on your responses. Please include a discussion regarding the sampling event in the report that is sent to DEQ with this form. For additional instructions, please click the Open Complete Instructions button.

[Open Complete Instructions](#)

**Basic Questions**

[View example \(Note: example optimized for viewing in Chrome browser\)](#)

1. Site/Facility name	<input type="text"/>	
2. Site code or facility ID (if applicable)	<input type="text"/>	
3. Release ID (if applicable)	<input type="text"/>	
4. Sample delivery group	<input type="text"/>	
5. Name of DEQ-approved sampling plan	<input type="text"/>	
6. Date DEQ approved the sampling plan	<input type="text"/>	M/D/YY
7. Name of data validator	<input type="text"/>	
8. Phone	<input type="text"/>	
9. Date validated	<input type="text"/>	M/D/YY

**Field Collection Questions**

[View example \(Note: example optimized for viewing in Chrome browser\)](#)

10. Sample matrix	<input type="checkbox"/> Soil	<input type="checkbox"/> Sediment	<input type="checkbox"/> Surface water	<input type="checkbox"/> Groundwater	<input type="checkbox"/> Tap water	<input type="checkbox"/> Air (including soil gas)	<input type="checkbox"/> Other <input type="text"/>
11. Sample collection start date	<input type="text"/>	M/D/YY					
12. Sample collection end date	<input type="text"/>	M/D/YY					
13. Analytical methods used <i>Use Add Method button to list multiple methods. Enter any other methods in the field manually.</i>	Add Method	<input type="text"/>					
	Delete Method	<input type="text"/>					

**Laboratory-related Questions**

[View example \(Note: example optimized for viewing in Chrome browser\)](#)

14. Laboratory name and location	<input type="text"/>			
15. Laboratory project ID	<input type="text"/>			
16. Were samples received in good condition and at appropriate temperature, chain-of-custody forms complete, and all samples analyzed within holding times?	Yes <input type="radio"/>	No <input type="radio"/>	See Below <input type="radio"/>	Comments <input type="text"/>
16a. Were chain-of-custody forms complete?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	

16b. Were samples received in good condition, preserved, and at appropriate temperature (VOA no headspace, appropriate pH, temperature 4° C +/- 2° for most samples)?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
16c. Were the samples analyzed within method-specified or technical holding times?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17. Were all laboratory quality control procedures complied with and is data validated without qualifiers?	Yes <input type="radio"/>	No <input type="radio"/>	See Below <input type="radio"/>	Comments <input type="text"/>
17a. Were all calibration verification results within acceptable limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17b. Were laboratory (method) blank samples free of contamination?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17c. Are the percent recoveries and relative percent differences of matrix spike and matrix spike duplicates within quality control limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17d. Are the laboratory control samples the same matrix as the samples and prepared the same as associated samples?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17e. Were laboratory control samples and laboratory control sample duplicate percent recoveries and relative percent differences within laboratory control limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17f. Were surrogate recoveries within laboratory quality control limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
17g. Were the laboratory duplicate relative percent differences within data validation quality control limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
18. Were the total number of lab method blanks at least 5% of the total number of samples, or as required by the method?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
19. Were the total number of lab matrix spike samples prepared at least 5% of the total number of samples, or as required by the method?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>	
20. Please list any project samples used for matrix spike/matrix spike duplicates.				
Add Sample	Lab ID	Field Sample ID	Comments	
Delete Sample				

21. Is the total number of laboratory control samples at least 5% of the total number of samples?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
---	------------------------------	-----------------------------	----------------------------------

**Consultant/Validator Questions**

[View example \(Note: example optimized for viewing in Chrome\)](#)

22. Are the detection limits appropriate for the project (i.e. at or below screening levels)?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
---	------------------------------	-----------------------------	----------------------------------

23. Are the reported units appropriate for the sample matrix (i.e. water results in ug/L, not mg/kg)?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
---	------------------------------	-----------------------------	----------------------------------

24. Do the analytical methods comply with project requirements (e.g. in the SAP, work plan, or QAPP)?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
---	------------------------------	-----------------------------	----------------------------------

25. Do the laboratory reports include all constituents requested to be analyzed on the chain-of-custody or under the sampling plan or other applicable document?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
--	------------------------------	-----------------------------	----------------------------------

26. Is the number of sample blanks (e.g. equipment, trip, or field blanks) equal to at least 10% of the total number of samples, or as otherwise required?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
--	------------------------------	-----------------------------	----------------------------------

27. Are field blanks free from contamination, duplicates collected as required, and field duplicate percent differences within data validation quality control limits?	Yes <input type="radio"/>	No <input type="radio"/>	See Below <input type="radio"/>	Comments <input type="text"/>
--	------------------------------	-----------------------------	------------------------------------	----------------------------------

27a. Were all blank samples free of analyte contamination?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
--	------------------------------	-----------------------------	----------------------------------

27b. Were field duplicates collected as required?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
---	------------------------------	-----------------------------	----------------------------------

27c. Are field duplicate relative percent differences within data validation quality control limits?	Yes <input type="radio"/>	No <input type="radio"/>	Comments <input type="text"/>
--	------------------------------	-----------------------------	----------------------------------

28. Please provide an Excel or CSV file to the DEQ project manager (via e-mail or CD) that lists all samples evaluated in this summary and lists any qualified data.   
Please use the following format:

Lab ID	Field Sample ID	Qualifiers	Comments (indicate whether the issue biases the results high or low)
Example 48310-2.31E	Example GW-1	R	Sample dropped in lab and unrecoverable
Example 48310-2.32D	Example GW-2		

Please use the following format for qualifiers. See EPA's National Functional Guidelines for more information on qualifiers for unique samples such as dioxins.

Qualifier	Explanation
C	Pesticide and Arochlor results confirmed with GC/MS
J-	Estimated value, may be biased low
J	Analyte identified, but concentration is estimated
J+	Estimated value, may be biased high
NJ	Tentatively identified compound
R	Sample result rejected
U	Analyte analyzed for, but not detected above quantitation limit
UJ	Analyte not detected above CRQL, but CRQL may be inaccurate
X	Pesticide and Arochlor results attempted using GC/MS, but unsuccessful

If you wish to manually enter qualified sample results, please use the table below.

Add Sample	Lab ID	Field Sample ID	Qualifiers	Comments (indicate whether the issue biases the results high or low)
Delete Sample				

29. What is the percent completeness (samples planned versus valid samples collected)? Comments

30. Was the completeness goal met? Comments

Yes  No

31. Does all data conform to analytical methods and data quality objectives specified for this project? Comments

Yes  No

32. Other general comments or observations?

**Split Samples**

33. Did DEQ collect split samples? Comments

Yes  No

Print Form

Save As

Open Instructions

Hide Instructions