



1435 Norjohn Court, Unit 1, Burlington, ON, Canada L7L 0E6

SVOC DATA PACKAGE

Client Project Information

Project ID: 1466-004 SEATTLE IRON & METALS

Project Description:

Contact: Amber Bailey

ALSE Project Information

Project ID: FAR100

Contact: Breanne Dusureault

Submission ID(s): L2541483

Final Package Review by:

A handwritten signature in black ink, appearing to read 'Breanne Dusureault', is written over a horizontal line.

Date Reviewed: 28-Jan-21

SVOC DATA PACKAGE

SECTION 1: PROJECT NARRATIVE

ALSE Project Information

 Project ID: FAR100
 Contact: Breanne Dusureault
 Submission ID(s): L2541483

Client Project Information

 Project ID: 1466-004 SEATTLE IRON & METALS
 Project Description:
 Contact: Amber Bailey

Analytical Method: PCDD/F by EPA TO9A/8290

| ALS Sample ID | Client Sample Descriptions | Matrix | Date Sampled | Date Received | Temp/degrees C. on receipt | Date Extracted | Date Analyzed |
|--------------------|--|----------------|--------------|---------------|----------------------------|------------------|------------------|
| L2519524-1 | L2497422-3-1 | PUF | 15-Oct-20 | 21-Oct-20 | 10.8 | n/a | n/a |
| L2530845-1 | L2497422-8-1 | PUF | 17-Nov-20 | 18-Nov-20 | 5.9 | n/a | n/a |
| L2541477-1 | L2527465-3-1 | PUF | 15-Dec-20 | 16-Dec-20 | 5.4 | n/a | n/a |
| L2541483-1 | SITE 1 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | PUF | n/a | n/a | n/a | 23-Dec-21 | 22-Jan-21 |
| L2519524-2 | L2497422-5-2 | PUF | 15-Oct-20 | 21-Oct-20 | 10.8 | n/a | n/a |
| L2530845-2 | L2516041-1-2 | PUF | 17-Nov-20 | 18-Nov-20 | 5.9 | n/a | n/a |
| L2541477-2 | L2527465-2-2 | PUF | 15-Dec-20 | 16-Dec-20 | 5.4 | n/a | n/a |
| L2541483-2 | SITE 2 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | PUF | n/a | n/a | n/a | 23-Dec-21 | 22-Jan-21 |
| L2519524-3 | L2497422-4-3 | PUF | 15-Oct-20 | 21-Oct-20 | 10.8 | n/a | n/a |
| L2530845-3 | L2497422-9-3 | PUF | 17-Nov-20 | 18-Nov-20 | 5.9 | n/a | n/a |
| L2541477-3 | L2527465-5-3 | PUF | 15-Dec-20 | 16-Dec-20 | 5.4 | n/a | n/a |
| L2541483-3 | SITE 3 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | PUF | n/a | n/a | n/a | 23-Dec-21 | 27-Jan-21 |
| L2519524-4 | L2497422-6-4 | PUF | 15-Oct-20 | 21-Oct-20 | 10.8 | n/a | n/a |
| L2530845-4 | L2497422-7-4 | PUF | 17-Nov-20 | 18-Nov-20 | 5.9 | n/a | n/a |
| L2541477-4 | L2527465-4-4 | PUF | 15-Dec-20 | 16-Dec-20 | 5.4 | n/a | n/a |
| L2541483-4 | SITE 4 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | PUF | n/a | n/a | n/a | 23-Dec-21 | 27-Jan-21 |
| L2519524-5 | L2497422-2-5 | PUF | 15-Oct-20 | 21-Oct-20 | 10.8 | n/a | n/a |
| L2530845-5 | L2497422-10-5 | PUF | 17-Nov-20 | 18-Nov-20 | 5.9 | n/a | n/a |
| L2541477-5 | L2527465-1-5 | PUF | 15-Dec-20 | 16-Dec-20 | 5.4 | n/a | n/a |
| L2541483-5 | SITE 5 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | PUF | n/a | n/a | n/a | 23-Dec-21 | 22-Jan-21 |
| WG3463499-1 | Method Blank | MEDIA | n/a | n/a | n/a | 23-Dec-21 | 27-Jan-21 |
| WG3463499-4 | Method Blank | REAGENT | n/a | n/a | n/a | 23-Dec-21 | 27-Jan-21 |
| WG3463499-2 | Laboratory Control Sample | MEDIA | n/a | n/a | n/a | 23-Dec-21 | 17-Jan-21 |

Comments and Notes:
a) Sample Integrity:

The samples were received on 3 different dates as noted above. The three samples for each sites were extracted together for a total of 5 composites. Some of the samples were received at above the recommended transportation and storage temperature. However, the brief period at above the recommended temperature is not expected to have a negative impact on reported native target results.

For the samples received 16-Dec-20, the samples were logged-in with identities supplied by the client, rather than the ones written on the chain-of-custody.

b) Instrumental Analysis:

The results for select samples have been reported from a reanalysis of the extract due to elevated instrument background.

No criteria failures or exceedances.

I certify that this data package is in compliance with the terms and condition of the contract, both technically and for completeness, for other than the conditions detailed above. Release of the data contained in this data package (hardcopy and/or electronic version) has been authorized by the Laboratory Manager or his designee, as verified by the following signature.



Steve Kennedy, Technical Supervisor

28-Jan-21

Date

SVOC DATA PACKAGE

SECTION 2: DATA SUMMARY REPORT



1435 Norjohn Court, Unit 1, Burlington, ON, Canada L7L 0E6
Phone: 905-331-3111, FAX: 905-331-4567

Certificate of Analysis

ALS Project Contact: Breanne Dusureault
ALS Project ID: FAR100
ALS WO#: L2541483
Date of Report: 29-Jan-21
Date of Sample Receipt: 16-Dec-20

Client Name: Farallon Consulting, L.L.C.
Client Address: 975 5th Avenue Northwest
Issaquah, WA 98027
USA
Client Contact: Amber Bailey
Client Project ID: 1466-004 SEATTLE IRON & METALS

COMMENTS: PCDD/F by EPA TO9A/8290

Certified by:

A handwritten signature in black ink, appearing to read "Steve Kennedy", is written over a horizontal line.

Steve Kennedy
Technical Supervisor

Results in this certificate relate only to the samples as submitted to the laboratory.

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Sample Analysis Summary Report

| Sample Name | SITE 1 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | SITE 2 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | SITE 3 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | SITE 4 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | SITE 5 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) |
|--|--|--|--|--|--|
| ALS Sample ID | L2541483-1 | L2541483-2 | L2541483-3 | L2541483-4 | L2541483-5 |
| Sample Size | 1 | 1 | 1 | 1 | 1 |
| Sample size units | SAMPLE | SAMPLE | SAMPLE | SAMPLE | SAMPLE |
| Percent Moisture | n/a | n/a | n/a | n/a | n/a |
| Sample Matrix | PUF | PUF | PUF | PUF | PUF |
| Sampling Date | n/a | n/a | n/a | n/a | n/a |
| Extraction Date | 23-Dec-21 | 23-Dec-21 | 23-Dec-21 | 23-Dec-21 | 23-Dec-21 |
| Target Analytes | pg | pg | pg | pg | pg |
| 2,3,7,8-TCDD | <2.1 | <1.5 | <3.1 | <3.4 | <1.5 |
| 1,2,3,7,8-PeCDD | 6.81 | <2.2 | 7.75 | <6.2 | 8.13 |
| 1,2,3,4,7,8-HxCDD | <6.0 | 2.92 | 8.00 | <4.9 | <9.4 |
| 1,2,3,6,7,8-HxCDD | <9.0 | <5.5 | <6.5 | 6.78 | 12.5 |
| 1,2,3,7,8,9-HxCDD | <11 | 4.18 | <6.7 | <13 | 16.0 |
| 1,2,3,4,6,7,8-HpCDD | 202 | 74.5 | 189 | 236 | 272 |
| OCDD | 1160 | 578 | 889 | 1040 | 1770 |
| 2,3,7,8-TCDF | <2.7 | 6.28 | <4.1 | <2.8 | 5.77 |
| 1,2,3,7,8-PeCDF | 2.99 | 4.29 | <4.6 | <4.1 | <3.1 |
| 2,3,4,7,8-PeCDF | 3.73 | 6.80 | 5.85 | <3.8 | 5.38 |
| 1,2,3,4,7,8-HxCDF | <3.1 | <4.5 | <3.4 | <4.3 | <5.0 |
| 1,2,3,6,7,8-HxCDF | 2.23 | 4.02 | <2.1 | <4.1 | <3.7 |
| 2,3,4,6,7,8-HxCDF | 3.15 | 4.85 | 2.38 | <4.5 | <2.7 |
| 1,2,3,7,8,9-HxCDF | <2.4 | <1.1 | <2.2 | <5.2 | <1.7 |
| 1,2,3,4,6,7,8-HpCDF | 17.6 | 22.7 | <18 | <28 | 27.5 |
| 1,2,3,4,7,8,9-HpCDF | 3.87 | <1.5 | <2.4 | <2.7 | <3.3 |
| OCDF | <24 | 27.2 | <21 | 27.0 | 51.4 |
| Field Spike Standards | % Rec | % Rec | % Rec | % Rec | % Rec |
| 37Cl4-2,3,7,8-TCDD | 98 | 97 | 93 | 95 | 97 |
| 13C12-1,2,3,4,7,8-HxCDD | 88 | 92 | 94 | 88 | 95 |
| 13C12-2,3,4,7,8-PeCDF | 109 | 107 | 116 | 113 | 113 |
| 13C12-1,2,3,4,7,8-HxCDF | 94 | 98 | 88 | 91 | 98 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 89 | 96 | 103 | 101 | 85 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 71 | 78 | 85 | 72 | 76 |
| 13C12-1,2,3,7,8-PeCDD | 64 | 73 | 86 | 68 | 62 |
| 13C12-1,2,3,6,7,8-HxCDD | 92 | 93 | 100 | 88 | 95 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 58 | 67 | 88 | 65 | 60 |
| 13C12-OCDD | 49 | 63 | 101 | 63 | 45 |
| 13C12-2,3,7,8-TCDF | 77 | 85 | 91 | 77 | 90 |
| 13C12-1,2,3,7,8-PeCDF | 69 | 79 | 85 | 70 | 69 |
| 13C12-1,2,3,6,7,8-HxCDF | 99 | 97 | 106 | 91 | 107 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 68 | 75 | 93 | 67 | 70 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 71 | 72 | 81 | 71 | 76 |
| Homologue Group Totals | pg | pg | pg | pg | pg |
| Total-TCDD | 13.4 | 14.5 | <3.1 | 18.7 | 14.8 |
| Total-PeCDD | 41.0 | 8.78 | 23.4 | 12.4 | 69.7 |
| Total-HxCDD | 59.1 | 39.3 | 51.4 | 67.3 | 90.6 |
| Total-HpCDD | 521 | 159 | 474 | 557 | 672 |
| Total-TCDF | 39.4 | 126 | 15.2 | 14.9 | 142 |
| Total-PeCDF | 16.3 | 43.0 | 14.0 | 8.47 | 68.2 |
| Total-HxCDF | 5.38 | 23.8 | 15.7 | <5.2 | 23.9 |
| Total-HpCDF | 21.4 | 22.7 | 10.7 | <2.7 | 53.2 |
| Toxic Equivalency - (WHO 2005) | | | | | |
| Lower Bound PCDD/F TEQ (WHO 2005) | 11.1 | 5.55 | 12.7 | 3.36 | 16.7 |
| Mid Point PCDD/F TEQ (WHO 2005) | 15.4 | 9.57 | 16.7 | 14.8 | 19.8 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 16.7 | 10.4 | 18.6 | 18.4 | 20.6 |

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Quality Control Summary Report

| Sample Name | Method Blank | Method Blank | Laboratory Control Sample |
|--|--------------|--------------|---------------------------|
| ALS Sample ID | WG3463499-1 | WG3463499-4 | WG3463499-2 |
| Sample Size | 1 | 1 | 1 |
| Sample size units | sample | sample | sample |
| Percent Moisture | n/a | n/a | n/a |
| Sample Matrix | MEDIA | REAGENT | MEDIA |
| Sampling Date | n/a | n/a | n/a |
| Extraction Date | 23-Dec-21 | 23-Dec-21 | 23-Dec-21 |
| Target Analytes | pg | pg | % Rec |
| 2,3,7,8-TCDD | <2.9 | <3.2 | 90 |
| 1,2,3,7,8-PeCDD | <1.7 | <1.7 | 105 |
| 1,2,3,4,7,8-HxCDD | <1.0 | <1.5 | 102 |
| 1,2,3,6,7,8-HxCDD | <0.90 | <1.3 | 100 |
| 1,2,3,7,8,9-HxCDD | <0.98 | <1.4 | 103 |
| 1,2,3,4,6,7,8-HpCDD | <2.5 | <2.2 | 98 |
| OCDD | 30.6 | <10 | 110 |
| 2,3,7,8-TCDF | <2.1 | <2.3 | 91 |
| 1,2,3,7,8-PeCDF | <1.0 | <1.0 | 90 |
| 2,3,4,7,8-PeCDF | <0.93 | <0.94 | 82 |
| 1,2,3,4,7,8-HxCDF | <0.85 | <0.93 | 90 |
| 1,2,3,6,7,8-HxCDF | <0.81 | <0.89 | 95 |
| 2,3,4,6,7,8-HxCDF | <0.89 | <0.98 | 86 |
| 1,2,3,7,8,9-HxCDF | <1.0 | <1.1 | 86 |
| 1,2,3,4,6,7,8-HpCDF | <2.2 | <0.85 | 89 |
| 1,2,3,4,7,8,9-HpCDF | <2.9 | <1.1 | 92 |
| OCDF | <3.0 | <3.7 | 79 |
| Field Spike Standards | % Rec | % Rec | % Rec |
| 37Cl4-2,3,7,8-TCDD | NS | NS | NS |
| 13C12-1,2,3,4,7,8-HxCDD | NS | NS | NS |
| 13C12-2,3,4,7,8-PeCDF | NS | NS | NS |
| 13C12-1,2,3,4,7,8-HxCDF | NS | NS | NS |
| 13C12-1,2,3,4,7,8,9-HpCDF | NS | NS | NS |
| Extraction Standards | | | |
| 13C12-2,3,7,8-TCDD | 68 | 53 | 52 |
| 13C12-1,2,3,7,8-PeCDD | 69 | 67 | 53 |
| 13C12-1,2,3,6,7,8-HxCDD | 68 | 67 | 56 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 73 | 72 | 54 |
| 13C12-OCDD | 73 | 69 | 59 |
| 13C12-2,3,7,8-TCDF | 70 | 60 | 56 |
| 13C12-1,2,3,7,8-PeCDF | 72 | 63 | 50 |
| 13C12-1,2,3,6,7,8-HxCDF | 72 | 71 | 56 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 77 | 78 | 53 |
| Cleanup Standard | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 70 | 61 | 54 |
| Homologue Group Totals | pg | pg | |
| Total-TCDD | <2.9 | <3.2 | |
| Total-PeCDD | <1.7 | <1.7 | |
| Total-HxCDD | <1.0 | <1.5 | |
| Total-HpCDD | <2.5 | <2.2 | |
| Total-TCDF | <2.1 | <2.3 | |
| Total-PeCDF | <1.0 | <1.0 | |
| Total-HxCDF | <1.0 | <1.1 | |
| Total-HpCDF | <2.9 | <1.1 | |
| Toxic Equivalency - (WHO 2005) | | | |
| Lower Bound PCDD/F TEQ (WHO 2005) | 0.00918 | 0.00 | |
| Mid Point PCDD/F TEQ (WHO 2005) | 2.93 | 3.15 | |
| Upper Bound PCDD/F TEQ (WHO 2005) | 5.85 | 6.30 | |

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Continuing Calibration Summary Report

| Sample Name | CVS | CCV | CCV | CCV | CCV | CCV | CCV |
|------------------------------|----------------|--------------|--------------|--------------|--------------|--------------|--------------|
| ALS Sample ID | H7-20-RS1-1209 | H7-21-CCV- | H7-21-CCV- | H7-21-CCV- | H7-21-CCV- | H7-21-CCV- | H7-21-CCV- |
| Sample Size | 1 | 1 | 1 | 1 | 1 | 1 | 1 |
| Sample size units | n/a | n/a | n/a | n/a | n/a | n/a | n/a |
| Percent Moisture | n/a | n/a | n/a | n/a | n/a | n/a | n/a |
| Sample Matrix | QC | QC | QC | QC | QC | QC | QC |
| Sampling Date | n/a | n/a | n/a | n/a | n/a | n/a | n/a |
| Extraction Date | n/a | n/a | n/a | n/a | n/a | n/a | n/a |
| Target Analytes | % Rec | % Rec | % Rec | % Rec | % Rec | % Rec | % Rec |
| 2,3,7,8-TCDD | 90 | 94 | 102 | 101 | 103 | 100 | 98 |
| 1,2,3,7,8-PeCDD | 106 | 103 | 105 | 109 | 110 | 104 | 106 |
| 1,2,3,4,7,8-HxCDD | 101 | 100 | 110 | 103 | 93 | 103 | 105 |
| 1,2,3,6,7,8-HxCDD | 94 | 97 | 99 | 118 | 104 | 109 | 113 |
| 1,2,3,7,8,9-HxCDD | 102 | 102 | 106 | 120 | 103 | 116 | 112 |
| 1,2,3,4,6,7,8-HpCDD | 98 | 100 | 104 | 111 | 112 | 107 | 105 |
| OCDD | 100 | 106 | 109 | 116 | 119 | 108 | 110 |
| 2,3,7,8-TCDF | 100 | 100 | 102 | 105 | 108 | 118 | 106 |
| 1,2,3,7,8-PeCDF | 106 | 89 | 91 | 98 | 97 | 95 | 99 |
| 2,3,4,7,8-PeCDF | 98 | 90 | 91 | 94 | 96 | 96 | 96 |
| 1,2,3,4,7,8-HxCDF | 100 | 89 | 94 | 96 | 96 | 101 | 102 |
| 1,2,3,6,7,8-HxCDF | 101 | 90 | 91 | 101 | 102 | 103 | 105 |
| 2,3,4,6,7,8-HxCDF | 100 | 90 | 94 | 101 | 102 | 107 | 107 |
| 1,2,3,7,8,9-HxCDF | 109 | 93 | 92 | 95 | 90 | 106 | 107 |
| 1,2,3,4,6,7,8-HpCDF | 101 | 93 | 98 | 107 | 108 | 104 | 103 |
| 1,2,3,4,7,8,9-HpCDF | 108 | 99 | 99 | 103 | 102 | 111 | 111 |
| OCDF | 102 | 85 | 88 | 101 | 104 | 100 | 102 |
| Field Spike Standards | % Rec | % Rec | % Rec | % Rec | % Rec | % Rec | % Rec |
| 37Cl4-2,3,7,8-TCDD | 107 | 90 | 91 | 91 | 92 | 91 | 90 |
| 13C12-1,2,3,4,7,8-HxCDD | 104 | 103 | 105 | 105 | 82 | 93 | 101 |
| 13C12-2,3,4,7,8-PeCDF | 100 | 101 | 99 | 98 | 99 | 98 | 99 |
| 13C12-1,2,3,4,7,8-HxCDF | 102 | 101 | 102 | 95 | 96 | 100 | 99 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 102 | 99 | 103 | 96 | 93 | 108 | 105 |
| Extraction Standards | | | | | | | |
| 13C12-2,3,7,8-TCDD | 101 | 104 | 105 | 104 | 104 | 103 | 106 |
| 13C12-1,2,3,7,8-PeCDD | 94 | 95 | 99 | 94 | 89 | 94 | 97 |
| 13C12-1,2,3,6,7,8-HxCDD | 94 | 96 | 96 | 95 | 107 | 99 | 98 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 102 | 104 | 107 | 99 | 94 | 115 | 116 |
| 13C12-OCDD | 105 | 127 | 120 | 109 | 101 | 165 | 150 |
| 13C12-2,3,7,8-TCDF | 97 | 97 | 105 | 108 | 109 | 101 | 105 |
| 13C12-1,2,3,7,8-PeCDF | 93 | 95 | 105 | 106 | 103 | 101 | 104 |
| 13C12-1,2,3,6,7,8-HxCDF | 93 | 92 | 99 | 108 | 111 | 103 | 106 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 95 | 94 | 101 | 102 | 99 | 113 | 115 |
| Cleanup Standard | | | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 95 | 96 | 101 | 104 | 99 | 108 | 111 |

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Sample Analysis Report

| | | | | |
|--------------------|---|------------------|-----------|---|
| Sample Name | SITE 1 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | Sampling Date | n/a | |
| ALS Sample ID | L2541483-1 | Extraction Date | 23-Dec-21 | Approved: <i>N Ashtari</i> --e-signature-- 28-Jan-2021 |
| Analysis Method | TO9A | Sample Size | 1 | |
| Analysis Type | Sample | Percent Moisture | n/a | |
| Sample Matrix | PUF | Split Ratio | 4 | |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210122A05 |
| Run Date | 22-Jan-21 12:43 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------------|------------|-----|
| 2,3,7,8-TCDD | 1 | 27.66 | <2.1 | 2.1 U | 2.0 | 20 |
| 1,2,3,7,8-PeCDD | 1 | 32.07 | 6.81 | 1.7 J | | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | 34.16 | <6.0 | 4.2 M,J,R | 6.0 | 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | 34.22 | <9.0 | 3.7 M,J,R | 9.0 | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | 34.35 | <11 | 4.0 M,J,R | 11 | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | 35.81 | 202 | 2.3 | | 100 |
| OCDD | 0.0003 | 37.26 | 1160 | 4.7 | | 200 |
| 2,3,7,8-TCDF | 0.1 | 26.68 | <2.7 | 2.7 M,U | 2.4 | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | 31.10 | 2.99 | 1.9 M,J | | 100 |
| 2,3,4,7,8-PeCDF | 0.3 | 31.76 | 3.73 | 1.8 M,J | | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | 33.67 | <3.1 | 2.0 M,J,R | 3.1 | 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | 33.73 | 2.23 | 1.9 J | | 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | 34.07 | 3.15 | 2.1 J | | 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | 34.49 | <2.4 | 2.4 M,U | 0.89 | 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | 35.25 | 17.6 | 0.90 J | | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | 36.05 | 3.87 | 1.2 J | | 100 |
| OCDF | 0.0003 | 37.36 | <24 | 4.2 J,R | 24 | 200 |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|-------|-------|------------|
| 37Cl4-2,3,7,8-TCDD | 1800 | 27.66 | 98 70-130 |
| 13C12-1,2,3,4,7,8-HxCDD | 18000 | 34.15 | 88 70-130 |
| 13C12-2,3,4,7,8-PeCDF | 18000 | 31.84 | 109 70-130 |
| 13C12-1,2,3,4,7,8-HxCDF | 18000 | 33.66 | 94 70-130 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 18000 | 36.05 | 89 70-130 |

| Extraction Standards | pg | Conc. | EDL |
|---------------------------|------|-------|-----------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.64 | 71 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.06 | 64 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.21 | 92 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.80 | 58 25-130 |
| 13C12-OCDD | 8000 | 37.26 | 49 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.72 | 77 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.07 | 69 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.72 | 99 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.25 | 68 25-130 |

| Cleanup Standard | pg | Conc. | EDL |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.48 | 71 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg | |
|------------------------|---------|-------------|-----------|-----|
| Total-TCDD | 3 | 13.4 | 2.1 | 20 |
| Total-PeCDD | 5 | 41.0 | 1.7 | 100 |
| Total-HxCDD | 2 | 59.1 | 4.2 | 100 |
| Total-HpCDD | 2 | 521 | 2.3 | 100 |
| Total-TCDF | 6 | 39.4 | 2.7 | 20 |
| Total-PeCDF | 4 | 16.3 | 1.9 | 100 |
| Total-HxCDF | 2 | 5.38 | 2.4 | 100 |
| Total-HpCDF | 2 | 21.4 | 1.2 | 100 |

| Toxic Equivalency - (WHO 2005) | pg |
|-----------------------------------|------|
| Lower Bound PCDD/F TEQ (WHO 2005) | 11.1 |
| Mid Point PCDD/F TEQ (WHO 2005) | 15.4 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 16.7 |

| | | |
|------|--|-------------------------------------|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. | |
| TEF | Indicates the Toxic Equivalency Factor | TEQ Indicates the Toxic Equivalency |
| M | Indicates that a peak has been manually integrated. | |
| U | Indicates that this compound was not detected above the EDL. | |
| J | Indicates that a target analyte was detected below the calibrated range. | |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. | |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. | |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure | |

ALS Life Sciences

Sample Analysis Report

| | | | | |
|--------------------|---|------------------|-----------|---|
| Sample Name | SITE 2 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | Sampling Date | n/a | |
| ALS Sample ID | L2541483-2 | Extraction Date | 23-Dec-21 | Approved: <i>N Ashtari</i> --e-signature-- 28-Jan-2021 |
| Analysis Method | TO9A | Sample Size | 1 | |
| Analysis Type | Sample | Percent Moisture | n/a | |
| Sample Matrix | PUF | Split Ratio | 4 | |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210122A06 |
| Run Date | 22-Jan-21 13:25 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg | Flags | EMPC pg | LQL |
|-------------------------------|-------------------|--------------|---------------------|-------------------|-------|------------|-----|
| 2,3,7,8-TCDD | 1 | NotFnd | <1.5 | 1.5 | U | | 20 |
| 1,2,3,7,8-PeCDD | 1 | 32.09 | <2.2 | 0.93 | M,J,R | 2.2 | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | 34.16 | 2.92 | 2.3 | J | | 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | 34.23 | <5.5 | 2.0 | J,R | 5.5 | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | 34.36 | 4.18 | 2.2 | J | | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | 35.82 | 74.5 | 1.3 | J | | 100 |
| OCDD | 0.0003 | 37.28 | 578 | 2.5 | | | 200 |
| 2,3,7,8-TCDF | 0.1 | 26.77 | 6.28 | 1.8 | M,J | | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | 31.11 | 4.29 | 1.9 | M,J | | 100 |
| 2,3,4,7,8-PeCDF | 0.3 | 31.85 | 6.80 | 1.7 | J | | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | 33.68 | <4.5 | 0.93 | J,R | 4.5 | 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | 33.74 | 4.02 | 0.89 | J | | 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | 34.08 | 4.85 | 0.98 | J | | 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | NotFnd | <1.1 | 1.1 | U | | 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | 35.26 | 22.7 | 1.0 | J | | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | 36.06 | <1.5 | 1.3 | M,J,R | 1.5 | 100 |
| OCDF | 0.0003 | 37.36 | 27.2 | 1.7 | J | | 200 |
| Field Spike Standards | pg | | % Rec | Limits | | | |
| 37Cl4-2,3,7,8-TCDD | 1800 | 27.67 | 97 | 70-130 | | | |
| 13C12-1,2,3,4,7,8-HxCDD | 18000 | 34.16 | 92 | 70-130 | | | |
| 13C12-2,3,4,7,8-PeCDF | 18000 | 31.85 | 107 | 70-130 | | | |
| 13C12-1,2,3,4,7,8-HxCDF | 18000 | 33.67 | 98 | 70-130 | | | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 18000 | 36.05 | 96 | 70-130 | | | |
| Extraction Standards | | | | | | | |
| 13C12-2,3,7,8-TCDD | 4000 | 27.64 | 78 | 40-130 | | | |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.07 | 73 | 40-130 | | | |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.22 | 93 | 40-130 | | | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.81 | 67 | 25-130 | | | |
| 13C12-OCDD | 8000 | 37.27 | 63 | 25-130 | | | |
| 13C12-2,3,7,8-TCDF | 4000 | 26.74 | 85 | 40-130 | | | |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.09 | 79 | 40-130 | | | |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.73 | 97 | 40-130 | | | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.26 | 75 | 25-130 | | | |
| Cleanup Standard | pg | | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.49 | 72 | 40-130 | | | |
| Homologue Group Totals | # peaks | | Conc. pg | EDL pg | | | |
| Total-TCDD | 2 | 14.5 | 1.5 | | | 20 | |
| Total-PeCDD | 2 | 8.78 | 0.93 | | | 100 | |
| Total-HxCDD | 5 | 39.3 | 2.3 | | | 100 | |
| Total-HpCDD | 2 | 159 | 1.3 | | | 100 | |
| Total-TCDF | 10 | 126 | 1.8 | | | 20 | |
| Total-PeCDF | 6 | 43.0 | 1.9 | | | 100 | |
| Total-HxCDF | 3 | 23.8 | 1.1 | | | 100 | |
| Total-HpCDF | 1 | 22.7 | 1.3 | | | 100 | |

| | |
|--|-----------|
| Toxic Equivalency - (WHO 2005) | pg |
| Lower Bound PCDD/F TEQ (WHO 2005) | 5.55 |
| Mid Point PCDD/F TEQ (WHO 2005) | 9.57 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 10.4 |

| | |
|------|--|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. |
| TEF | Indicates the Toxic Equivalency Factor |
| M | Indicates that a peak has been manually integrated. |
| U | Indicates that this compound was not detected above the EDL. |
| J | Indicates that a target analyte was detected below the calibrated range. |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure |

ALS Life Sciences

Sample Analysis Report

| | | | |
|--|------------------|-----------|--------|
| Sample Name SITE 3 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | Sampling Date | n/a | |
| ALS Sample ID L2541483-3 | Extraction Date | 23-Dec-21 | |
| Analysis Method TO9A | Sample Size | 1 | Sample |
| Analysis Type Sample | Percent Moisture | n/a | |
| Sample Matrix PUF | Split Ratio | 4 | |

Approved:
N Ashtari
--e-signature--
28-Jan-2021

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A09 |
| Run Date | 27-Jan-21 22:40 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg | Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------|-------|------------|-----|
| 2,3,7,8-TCDD | 1 | NotFnd | <3.1 | 3.1 | U | | 20 |
| 1,2,3,7,8-PeCDD | 1 | 32.06 | 7.75 | 2.4 | J | | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | 34.15 | 8.00 | 2.2 | M,J | | 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | 34.20 | <6.5 | 1.9 | M,J,R | 6.5 | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | 34.32 | <6.7 | 2.1 | M,J,R | 6.7 | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | 35.79 | 189 | 4.2 | | | 100 |
| OCDD | 0.0003 | 37.24 | 889 | 5.7 | | | 200 |
| 2,3,7,8-TCDF | 0.1 | 26.75 | <4.1 | 4.1 | M,U | 2.2 | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | NotFnd | <4.6 | 4.6 | U | | 100 |
| 2,3,4,7,8-PeCDF | 0.3 | 31.75 | 5.85 | 4.2 | J | | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | 33.66 | <3.4 | 1.8 | M,J,R | 3.4 | 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | 33.72 | <2.1 | 1.8 | M,J,R | 2.1 | 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | 34.06 | 2.38 | 1.9 | M,J | | 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | 34.47 | <2.2 | 2.2 | M,U | | 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | 35.24 | <18 | 1.8 | M,J,R | 18 | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | 36.04 | <2.4 | 2.4 | M,U | 2.2 | 100 |
| OCDF | 0.0003 | 37.32 | <21 | 1.7 | M,J,R | 21 | 200 |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|-------|-------|------------|
| 37Cl4-2,3,7,8-TCDD | 1800 | 27.66 | 93 70-130 |
| 13C12-1,2,3,4,7,8-HxCDD | 18000 | 34.14 | 94 70-130 |
| 13C12-2,3,4,7,8-PeCDF | 18000 | 31.83 | 116 70-130 |
| 13C12-1,2,3,4,7,8-HxCDF | 18000 | 33.65 | 88 70-130 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 18000 | 36.03 | 103 70-130 |

| Extraction Standards | pg | Conc. | EDL |
|---------------------------|------|-------|------------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.63 | 85 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.05 | 86 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.20 | 100 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.79 | 88 25-130 |
| 13C12-OCDD | 8000 | 37.24 | 101 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.72 | 91 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.07 | 85 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.72 | 106 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.23 | 93 25-130 |

| Cleanup Standard | pg | Conc. | EDL |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.47 | 81 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg |
|------------------------|---------|-------------|-----------|
| Total-TCDD | 0 | <3.1 | 3.1 |
| Total-PeCDD | 3 | 23.4 | 2.4 |
| Total-HxCDD | 2 | 51.4 | 2.2 |
| Total-HpCDD | 2 | 474 | 4.2 |
| Total-TCDF | 3 | 15.2 | 4.1 |
| Total-PeCDF | 3 | 14.0 | 4.6 |
| Total-HxCDF | 3 | 15.7 | 2.2 |
| Total-HpCDF | 1 | 10.7 | 2.4 |

| | |
|--|-----------|
| Toxic Equivalency - (WHO 2005) | pg |
| Lower Bound PCDD/F TEQ (WHO 2005) | 12.7 |
| Mid Point PCDD/F TEQ (WHO 2005) | 16.7 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 18.6 |

| | |
|------|--|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. |
| TEF | Indicates the Toxic Equivalency Factor |
| M | Indicates that a peak has been manually integrated. |
| U | Indicates that this compound was not detected above the EDL. |
| J | Indicates that a target analyte was detected below the calibrated range. |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure |

ALS Life Sciences

Sample Analysis Report

| | | | |
|--|------------------|-----------|--------|
| Sample Name SITE 4 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | Sampling Date | n/a | |
| ALS Sample ID L2541483-4 | Extraction Date | 23-Dec-21 | |
| Analysis Method TO9A | Sample Size | 1 | Sample |
| Analysis Type Sample | Percent Moisture | n/a | |
| Sample Matrix PUF | Split Ratio | 4 | |

Approved:
N Ashtari
--e-signature--
28-Jan-2021

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A10 |
| Run Date | 27-Jan-21 23:22 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg | Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------|-------|------------|-----|
| 2,3,7,8-TCDD | 1 | NotFnd | <3.4 | 3.4 | | U | 20 |
| 1,2,3,7,8-PeCDD | 1 | 32.07 | <6.2 | 3.1 | M,J,R | 6.2 | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | 34.15 | <4.9 | 4.9 | M,U | 3.8 | 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | 34.20 | 6.78 | 4.3 | M,J | | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | 34.33 | <13 | 4.7 | M,J,R | 13 | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | 35.79 | 236 | 7.5 | | | 100 |
| OCDD | 0.0003 | 37.24 | 1040 | 8.1 | | | 200 |
| 2,3,7,8-TCDF | 0.1 | NotFnd | <2.8 | 2.8 | | U | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | NotFnd | <4.1 | 4.1 | | U | 100 |
| 2,3,4,7,8-PeCDF | 0.3 | 31.84 | <3.8 | 3.8 | M,U | 2.7 | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | 33.66 | <4.3 | 4.3 | M,U | 2.6 | 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | NotFnd | <4.1 | 4.1 | | U | 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | 34.06 | <4.5 | 4.5 | M,U | 4.4 | 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | 34.49 | <5.2 | 5.2 | M,U | 1.1 | 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | 35.24 | <28 | 2.1 | J,R | 28 | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | NotFnd | <2.7 | 2.7 | | U | 100 |
| OCDF | 0.0003 | 37.33 | 27.0 | 14 | J | | 200 |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|-------|-------|------------|
| 37C14-2,3,7,8-TCDD | 1800 | 27.66 | 95 70-130 |
| 13C12-1,2,3,4,7,8-HxCDD | 18000 | 34.14 | 88 70-130 |
| 13C12-2,3,4,7,8-PeCDF | 18000 | 31.83 | 113 70-130 |
| 13C12-1,2,3,4,7,8-HxCDF | 18000 | 33.65 | 91 70-130 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 18000 | 36.03 | 101 70-130 |

| Extraction Standards | pg | % Rec | Limits |
|---------------------------|------|-------|-----------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.63 | 72 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.05 | 68 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.20 | 88 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.79 | 65 25-130 |
| 13C12-OCDD | 8000 | 37.23 | 63 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.72 | 77 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.06 | 70 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.71 | 91 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.23 | 67 25-130 |

| Cleanup Standard | pg | % Rec | Limits |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HpCDF | 4000 | 34.47 | 71 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg | | |
|------------------------|---------|-------------|-----------|-----|-----|
| Total-TCDD | 3 | 18.7 | 3.4 | 20 | |
| Total-PeCDD | 1 | 12.4 | 3.1 | 100 | |
| Total-HxCDD | 2 | 67.3 | 4.9 | 100 | |
| Total-HpCDD | 2 | 557 | 7.5 | 100 | |
| Total-TCDF | 4 | 14.9 | 2.8 | 20 | |
| Total-PeCDF | 2 | 8.47 | 4.1 | 100 | |
| Total-HxCDF | 0 | <5.2 | 5.2 | U | 100 |
| Total-HpCDF | 0 | <2.7 | 2.7 | U | 100 |

| | |
|--|-----------|
| Toxic Equivalency - (WHO 2005) | pg |
| Lower Bound PCDD/F TEQ (WHO 2005) | 3.36 |
| Mid Point PCDD/F TEQ (WHO 2005) | 14.8 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 18.4 |

| | |
|------|--|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. |
| TEF | Indicates the Toxic Equivalency Factor |
| M | Indicates that a peak has been manually integrated. |
| U | Indicates that this compound was not detected above the EDL. |
| J | Indicates that a target analyte was detected below the calibrated range. |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure |

ALS Life Sciences

Sample Analysis Report

| | | | | |
|--------------------|---|------------------|-----------|---|
| Sample Name | SITE 5 - COMPOSITE 4 (WET SEASON - OCT, NOV, DEC) | Sampling Date | n/a | |
| ALS Sample ID | L2541483-5 | Extraction Date | 23-Dec-21 | Approved: <i>N Ashtari</i> --e-signature-- 28-Jan-2021 |
| Analysis Method | TO9A | Sample Size | 1 Sample | |
| Analysis Type | Sample | Percent Moisture | n/a | |
| Sample Matrix | PUF | Split Ratio | 4 | |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210122A07 |
| Run Date | 22-Jan-21 14:07 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------------|------------|---------|
| 2,3,7,8-TCDD | 1 | NotFnd | <1.5 | 1.5 | U | 20 |
| 1,2,3,7,8-PeCDD | 1 | 32.08 | 8.13 | 1.2 | J | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | 34.18 | <9.4 | 2.8 | M,J,R | 9.4 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | 34.23 | 12.5 | 2.5 | M,J | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | 34.35 | 16.0 | 2.7 | M,J | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | 35.82 | 272 | 2.6 | | 100 |
| OCDD | 0.0003 | 37.27 | 1770 | 7.0 | | 200 |
| 2,3,7,8-TCDF | 0.1 | 26.75 | 5.77 | 1.1 | M,J | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | 31.11 | <3.1 | 2.7 | M,J,R | 3.1 100 |
| 2,3,4,7,8-PeCDF | 0.3 | 31.85 | 5.38 | 2.5 | M,J | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | 33.68 | <5.0 | 1.4 | M,J,R | 5.0 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | 33.74 | <3.7 | 1.3 | J,R | 3.7 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | 34.08 | <2.7 | 1.5 | J,R | 2.7 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | 34.50 | <1.7 | 1.7 | M,U | 1.5 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | 35.26 | 27.5 | 1.1 | J | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | 36.08 | <3.3 | 1.4 | M,J,R | 3.3 100 |
| OCDF | 0.0003 | 37.36 | 51.4 | 2.2 | J | 200 |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|-------|-------|------------|
| 37Cl4-2,3,7,8-TCDD | 1800 | 27.67 | 97 70-130 |
| 13C12-1,2,3,4,7,8-HxCDD | 18000 | 34.16 | 95 70-130 |
| 13C12-2,3,4,7,8-PeCDF | 18000 | 31.85 | 113 70-130 |
| 13C12-1,2,3,4,7,8-HxCDF | 18000 | 33.67 | 98 70-130 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 18000 | 36.05 | 85 70-130 |

| Extraction Standards | pg | % Rec | Limits |
|---------------------------|------|-------|------------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.64 | 76 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.07 | 62 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.22 | 95 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.81 | 60 25-130 |
| 13C12-OCDD | 8000 | 37.27 | 45 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.74 | 90 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.09 | 69 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.73 | 107 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.25 | 70 25-130 |

| Cleanup Standard | pg | % Rec | Limits |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.49 | 76 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg | |
|------------------------|---------|-------------|-----------|-----|
| Total-TCDD | 3 | 14.8 | 1.5 | 20 |
| Total-PeCDD | 6 | 69.7 | 1.2 | 100 |
| Total-HxCDD | 4 | 90.6 | 2.8 | 100 |
| Total-HpCDD | 2 | 672 | 2.6 | 100 |
| Total-TCDF | 12 | 142 | 1.1 | 20 |
| Total-PeCDF | 8 | 68.2 | 2.7 | 100 |
| Total-HxCDF | 2 | 23.9 | 1.7 | 100 |
| Total-HpCDF | 2 | 53.2 | 1.4 | 100 |

| Toxic Equivalency - (WHO 2005) | pg |
|-----------------------------------|------|
| Lower Bound PCDD/F TEQ (WHO 2005) | 16.7 |
| Mid Point PCDD/F TEQ (WHO 2005) | 19.8 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 20.6 |

EDL Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample.
 TEF Indicates the Toxic Equivalency Factor TEQ Indicates the Toxic Equivalency
 M Indicates that a peak has been manually integrated.
 U Indicates that this compound was not detected above the EDL.
 J Indicates that a target analyte was detected below the calibrated range.
 R Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion.
 LQL Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions.
 EMPC Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure

SVOC DATA PACKAGE

SECTION 3: METHOD SUMMARY

PCDD/F METHOD SUMMARY
Methods 23/0023A/1613B/8290/TO-9A

Introduction:

This summary is to provide ALSE Burlington PCDD/F method details in order to provide persons reviewing or validating this data package sufficient information to re-construct the sample calculation, data verification and review. It incorporates the analysis of PCDD/F via the following reference methods:

- US EPA Office of Water, Method 1613B
- US EPA Office of Solid Waste, SW846 Methods 8290A and 0023/8290A
- US EPA Office of Research & Development Method TO-9A.
- US EPA Office of Air Quality Planning & Standards Method 23.

Any deviations to what is listed herein would be listed in the project narrative.

To avoid the confusion and conflicting nomenclature within the methods, we have defined the labeled standards in terms relating to the time of addition to the sample or extract. Therefore;

- The Field or Sampling Standards are added prior to field sampling
- The Extraction Standards are added prior to extraction
- The Clean-up Standards are added prior to extract clean-up
- The Injection Standards are added prior to extract injection.

Calibration Standard Levels:

Six levels of standard are available for calibration as listed in Table 1. The low point (the CS0) is below method requirements and therefore is optional.

Table 1: Calibration Standards

| | CS0 | CS1 | CS2 | CS3 | CS4 | CS5 | |
|---|--|------------|------------|------------|------------|------------|------|
| Natives | 2,3,7,8-TCDD | 0.1 | 0.5 | 2 | 10 | 40 | 200 |
| | 2,3,7,8-TCDF | 0.1 | 0.5 | 2 | 10 | 40 | 200 |
| | 1,2,3,7,8-PeCDD | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,7,8-PeCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 2,3,4,7,8-PeCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,4,7,8-HxCDD | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,6,7,8-HxCDD | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,7,8,9-HxCDD | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,4,7,8-HxCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,6,7,8-HxCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,7,8,9-HxCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 2,3,4,6,7,8-HxCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,4,6,7,8-HpCDD | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,4,6,7,8-HpCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | 1,2,3,4,7,8,9-HpCDF | 0.5 | 2.5 | 10 | 50 | 200 | 1000 |
| | OCDD | 1 | 5 | 20 | 100 | 400 | 2000 |
| | OCDF | 1 | 5 | 20 | 100 | 400 | 2000 |
| Labeled | 2,3,7,8-TCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 2,3,7,8-TCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | OCDD- ¹³ C ₁₂ | 200 | 200 | 200 | 200 | 200 | 200 |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | 0.1 | 0.5 | 2 | 10 | 40 | 200 | |
| Injection | 1,2,3,4-TCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |
| | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 100 | 100 | 100 | 100 | 100 | 100 |

Calibration Control Limits

The initial and continuing calibration control limits for all methods are presented in Table 2 below. For the initial calibration CS1 and for each calibration verification CS3, the signal to noise ratio for each quantification ion for labelled and non-labelled analytes must be greater than or equal to 10:1

Table 2: Calibration Control Limits

| | 1613B | | M23 & TO-9A | | 8290A | |
|--|--------------|-----------|----------------|-----------|--------------|-----------|
| | Initial Cal. | Cal. Ver. | Initial Cal. | Cal. Ver. | Initial Cal. | Cal. Ver. |
| | %RSD | ng/mL | %RSD | % Diff | %RSD | % Diff |
| Natives | | | | | | |
| 2,3,7,8-TCDD | 20 | 7.8-12.9 | 25 | 25 | 20 | 20* |
| 2,3,7,8-TCDF | 20 | 8.4-12.0 | 25 | 25 | 20 | 20* |
| 1,2,3,7,8-PeCDD | 20 | 39-65 | 25 | 25 | 20 | 20* |
| 1,2,3,7,8-PeCDF | 20 | 41-60 | 25 | 25 | 20 | 20* |
| 2,3,4,7,8-PeCDF | 20 | 41-61 | 25 | 25 | 20 | 20* |
| 1,2,3,4,7,8-HxCDD | 20 | 39-64 | 25 | 25 | 20 | 20* |
| 1,2,3,6,7,8-HxCDD | 20 | 39-64 | 25 | 25 | 20 | 20* |
| 1,2,3,7,8,9-HxCDD | 35 | 41-61 | 25 | 25 | 20 | 20* |
| 1,2,3,4,7,8-HxCDF | 20 | 45-56 | 25 | 25 | 20 | 20* |
| 1,2,3,6,7,8-HxCDF | 20 | 44-57 | 25 | 25 | 20 | 20* |
| 1,2,3,7,8,9-HxCDF | 20 | 45-56 | 25 | 25 | 20 | 20* |
| 2,3,4,6,7,8-HxCDF | 20 | 44-57 | 25 | 25 | 20 | 20* |
| 1,2,3,4,6,7,8-HpCDD | 20 | 43-58 | 25 | 25 | 20 | 20* |
| 1,2,3,4,6,7,8-HpCDF | 20 | 45-55 | 25 | 25 | 20 | 20* |
| 1,2,3,4,7,8,9-HpCDF | 20 | 43-58 | 25 | 25 | 20 | 20* |
| OCDD | 20 | 79-126 | 25 | 25 | 20 | 20* |
| OCDF | 35 | 63-159 | 30 | 30 | 20 | 20* |
| Labels | | | | | | |
| 2,3,7,8-TCDD- ¹³ C ₁₂ | 35 | 82-121 | 25 | 25 | 30 | 30** |
| 2,3,7,8-TCDF- ¹³ C ₁₂ | 35 | 71-140 | 30 | 30 | 30 | 30** |
| 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 35 | 62-160 | 30 | 30 | 30 | 30** |
| 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 35 | 76-130 | 30 | 30 | 30 | 30** |
| 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 35 | 77-130 | 25 | 25 | 30 | 30** |
| 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 35 | 85-117 | 25 | 25 | 30 | 30** |
| 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 35 | 85-118 | 25 | 25 | 30 | 30** |
| 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 35 | 76-131 | 25 | 25 | 30 | 30** |
| 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 35 | 70-143 | 30 | 30 | 30 | 30** |
| 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 35 | 74-135 | - | - | - | - |
| 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 35 | 73-137 | 30 | 30 | 30 | 30** |
| 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 35 | 72-138 | 30 | 30 | 30 | 30** |
| 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 35 | 78-129 | 30 | 30 | 30 | 30** |
| 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 35 | 77-129 | 25 | 25 | 30 | 30** |
| OCDD- ¹³ C ₁₂ | 35 | 96-415 | 30 | 30 | 30 | 30** |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | 35 | 7.9-12.7 | 25 | 25 | 30 | 30** |

* 25% is allowed for a post-run verification but when the value is above 20%, then the analyte quantification must be as per 8290A Section 8.3.2.4 and corrective action is required before more samples can be analyzed.

**35% is allowed for a post-run verification but when the value is above 30%, then the analyte quantification must be as per 8290A Section 8.3.2.4 and corrective action is required before more samples can be analyzed.

LCS Criteria:

The laboratory control sample (LCS) or the On-Going Precision and Accuracy (OPR) recovery criteria are listed in Table 3

Table 3: Acceptance Criteria for IPR and OPR^a

| | Test Conc. | IPR | | OPR |
|--|---------------|----------------|----------------|----------|
| | | s ^b | X ^c | |
| | ng/L | ng/L | ng/L | ng/L |
| Natives | | | | |
| 2,3,7,8-TCDD | 10 | 2.8 | 8.3-12.9 | 6.7-15.8 |
| 2,3,7,8-TCDF | 10 | 2 | 8.7-13.7 | 7.5-15.8 |
| 1,2,3,7,8-PeCDD | 50 | 7.5 | 38-66 | 35-71 |
| 1,2,3,7,8-PeCDF | 50 | 7.5 | 43-62 | 40-67 |
| 2,3,4,7,8-PeCDF | 50 | 8.6 | 36-75 | 34-80 |
| 1,2,3,4,7,8-HxCDD | 50 | 9.4 | 39-76 | 35-82 |
| 1,2,3,6,7,8-HxCDD | 50 | 7.7 | 42-62 | 38-67 |
| 1,2,3,7,8,9-HxCDD | 50 | 11.1 | 37-71 | 32-81 |
| 1,2,3,4,7,8-HxCDF | 50 | 8.7 | 41-59 | 36-67 |
| 1,2,3,6,7,8-HxCDF | 50 | 6.7 | 46-60 | 42-65 |
| 1,2,3,7,8,9-HxCDF | 50 | 6.4 | 42-61 | 39-65 |
| 2,3,4,6,7,8-HxCDF | 50 | 7.4 | 37-74 | 35-78 |
| 1,2,3,4,6,7,8-HpCDD | 50 | 7.7 | 38-65 | 35-70 |
| 1,2,3,4,6,7,8-HpCDF | 50 | 6.3 | 45-56 | 41-61 |
| 1,2,3,4,7,8,9-HpCDF | 50 | 8.1 | 43-63 | 39-69 |
| OCDD | 100 | 19 | 89-127 | 78-144 |
| OCDF | 100 | 27 | 74-146 | 63-170 |
| Labels | | | | |
| 2,3,7,8-TCDD- ¹³ C ₁₂ | 100 | 37 | 28-134 | 20-175 |
| 2,3,7,8-TCDF- ¹³ C ₁₂ | 100 | 35 | 31-113 | 22-152 |
| 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 100 | 39 | 27-184 | 21-227 |
| 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 100 | 34 | 27-156 | 21-192 |
| 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 100 | 38 | 16-297 | 13-328 |
| 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 100 | 41 | 29-147 | 21-193 |
| 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 100 | 38 | 34-122 | 25-163 |
| 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 100 | 43 | 27-152 | 19-202 |
| 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 100 | 35 | 30-122 | 21-159 |
| 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 100 | 40 | 24-157 | 17-205 |
| 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 100 | 37 | 29-136 | 22-176 |
| 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 100 | 35 | 34-129 | 26-166 |
| 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 100 | 41 | 32-110 | 21-158 |
| 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 100 | 40 | 28-141 | 20-186 |
| OCDD- ¹³ C ₁₂ | 200 | 95 | 41-276 | 26-397 |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | 10 | 3.6 | 3.9-15.4 | 3.1-19.1 |

^a Assuming a final volume of 20uL

^b s = standard deviation

^c X = Average Concentration

Extraction/Clean-up & Sampling Standard Recovery Limits:

Table 4: Extraction, Clean-up, Injection & Sampling Standard Recovery Limits

| | 1613B or 8290A (non Stack) | | M23 or 0023A/8290A or TO-9A | |
|--|-------------------------------|------|-----------------------------------|------|
| | (% Rec.) | Ref. | (% Rec.) | Ref. |
| Extraction Standard | | | | |
| 2,3,7,8-TCDD- ¹³ C ₁₂ | 25-164 | a | 40-130 | b |
| 2,3,7,8-TCDF- ¹³ C ₁₂ | 24-169 | a | 40-130 | b |
| 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 25-181 | a | 40-130 | b |
| 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 24-185 | a | 40-130 | b |
| 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 21-178 | a | - | |
| 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 32-141 | a | - | |
| 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 28-130 | a | 40-130 | b |
| 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 26-152 | a | - | |
| 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 26-123 | a | 40-130 | b |
| 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 29-147 | a | - | |
| 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 28-136 | a | 40-130 | c,d |
| 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 23-140 | a | 25-130 | b |
| 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 28-143 | a | 25-130 | b |
| 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 26-138 | a | - | |
| OCDD- ¹³ C ₁₂ | 17-157 | a | 25-130 | b |
| Clean-up Standard | | | | |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | 35-197 | a | - | |
| 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | - | | 40-130 | b |
| Injection Standard | | | | |
| 1,2,3,4-TCDD- ¹³ C ₁₂ | 30-300 | d | 30-300 | d |
| 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 30-300 | d | 30-300 | d |
| Sampling Standard | | | | |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | - | | 70-130 | b |
| 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | - | | 70-130 | b |
| 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | - | | 70-130 | b |
| 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | - | | 70-130 | b |
| 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | - | | 70-130 | b |

References & Notes

^a from OW method 1613B

^b from OAQPS method 23

^c this extraction standard is not required in methods 23 and 0023A/8290A

^d ALS In-house criteria

Reporting Limits:

Unless indicated in the otherwise, the PCDD/F data is reported down to 2.5:1 signal to noise for each isomer grouping for each extract injection. This is consistent to SW846 8290 defined protocols (i.e. EDL or Estimated Detection Limit) and is commonly applied throughout the industry to all the HRMS PCDD/F methods applicable to this method summary.

Method Blank:

The method blank levels must be below the response to the low calibration standard, CS0 or CS1, whichever low calibration point is being applied to the project.

MS/MSD:

The % relative difference between the MS and MSD spike recoveries should be less than or equal to 20%.

Instrument/Run Performance Criteria:

- 1 Elution windows must be defined by a 'Window Performance Mix' at the beginning of each 12-hour run sequence
- 2 GC performance criteria of 25% maximum valley between 2,3,7,8-TCDD and its nearest eluting isomers (DB5) or 2,3,7,8-TCDF and its nearest eluting isomers (DB225).
- 3 At the beginning of and just following the end of each 12 hour run sequence, the instrument must be checked to demonstrate a resolution of 10,000 for each quantification window.
- 4 For method 1613B, the relative retention times (RRT) of the compounds in the daily CS3 calibration verification must fall into the ranges presented in Table 4.
- 5 For all calibrations, QC samples and field samples, the absolute retention time (RT) for 1,2,3,4-TCDD-13C12 must be >25.0 min on a DB5 column and >15.0 min on a DB225 column.
- 6 The RT in the daily CS3 verification standards must be within 15 seconds of the CS3 in the initial calibration run.
- 7 The maximum time between scans within a descriptor is 1 second.
- 8 Lock mass deviations to the average response must be less than or equal 20%.

Laboratory Duplicates:

The % relative difference between duplicates should be less than or equal to 25% but only where the response is greater than the low calibration standard.

Analyte Identification Criteria:

- 1 Ion ratio must be within 15% of theoretical or within 10% of the most recent CS3.
- 2 The retention time (RT) of the peak maxima for each pair of quantification ions must be no more than 2 seconds (i.e. 2 scans) difference.
- 3 The retention time (RT) of the peak maxima of all 2,3,7,8- substituted native analytes must be within -1 to +3 seconds of the RT of corresponding ¹³C₁₂-labelled isomer of that injection run.
- 4 For those native analytes without a corresponding labelled isomer, the relative retention time (RRT) must be within 0.005 of the relative retention time observed in the daily CS3 run.
- 5 When there is a significant PCDPE interference observed, then a peak in the PCDF channel is not confirmed to be PCDF. [Significant PCDPE interference is identified when there is a PCDPE parent ion peak 10% or more of the response of a peak at the same RT (i.e. within 2 seconds) in the corresponding PCDF channel.]
- 6 For any peak to be identified as a positive PCDD/F response, that peak must be within the retention time windows defined by the daily analysis of Window Performance Mixture.

Table 4: Quantitation References and Method 1613B RT References and RRT

| Analyte | Stack/Ambient Quantitation Reference | Method 1613B RT Reference | Method 1613B RRT |
|--|--|--|------------------|
| | | Solids/ Waters Quantitation Reference | |
| Compounds using 1,2,3,4-TCDD-¹³C₁₂ as injection standard | | | |
| 2,3,7,8-TCDF | 2,3,7,8-TCDF- ¹³ C ₁₂ | 2,3,7,8-TCDF- ¹³ C ₁₂ | 0.999-1.003 |
| 2,3,7,8-TCDD | 2,3,7,8-TCDD- ¹³ C ₁₂ | 2,3,7,8-TCDD- ¹³ C ₁₂ | 0.999-1.002 |
| 1,2,3,7,8-PeCDF | 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 0.999-1.002 |
| 2,3,4,7,8-PeCDF | 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 0.999-1.002 |
| 1,2,3,7,8-PeCDD | 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 0.999-1.002 |
| 2,3,7,8-TCDF- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 0.923-1.103 |
| 2,3,7,8-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 0.976-1.043 |
| 2,3,7,8-TCDD- ³⁷ Cl ₄ | 2,3,7,8-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 0.989-1.052 |
| 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1.000-1.425 |
| 2,3,4,7,8-PeCDF- ¹³ C ₁₂ | 1,2,3,7,8-PeCDF- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1.011-1.526 |
| 1,2,3,7,8-PeCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1,2,3,4-TCDD- ¹³ C ₁₂ | 1.000-1.567 |
| Compounds using 1,2,3,7,8,9-HxCDD-¹³C₁₂ as injection standard | | | |
| 1,2,3,4,7,8-HxCDF | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,6,7,8-HxCDF | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 0.997-1.005 |
| 1,2,3,7,8,9-HxCDF | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 0.999-1.001 |
| 2,3,4,6,7,8-HxCDF | 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,4,7,8-HxCDD | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,6,7,8-HxCDD | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 0.998-1.004 |
| 1,2,3,7,8,9-HxCDD ^a | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | ^a | 1.000-1.019 |
| 1,2,3,4,6,7,8-HpCDF | 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,4,7,8,9-HpCDF | 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,4,6,7,8-HpCDD | 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 0.999-1.001 |
| OCDF | OCDD- ¹³ C ₁₂ | OCDD- ¹³ C ₁₂ | 0.999-1.008 |
| OCDD | OCDD- ¹³ C ₁₂ | OCDD- ¹³ C ₁₂ | 0.999-1.001 |
| 1,2,3,4,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.944-0.970 |
| 1,2,3,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.949-0.975 |
| 1,2,3,7,8,9-HxCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.977-1.047 |
| 2,3,4,6,7,8-HxCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.959-1.021 |
| 1,2,3,4,7,8-HxCDD- ¹³ C ₁₂ | 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.977-1.000 |
| 1,2,3,6,7,8-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 0.981-1.003 |
| 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1.043-1.085 |
| 1,2,3,4,7,8,9-HpCDF- ¹³ C ₁₂ | 1,2,3,4,6,7,8-HpCDF- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1.057-1.151 |
| 1,2,3,4,6,7,8-HpCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1.086-1.110 |
| OCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1,2,3,7,8,9-HxCDD- ¹³ C ₁₂ | 1.032-1.311 |

^a For solids/waters via 1612B, 1,2,3,7,8,9-HxCDD is quantified against the average responses of 1,2,3,4,7,8-HxCDD-¹³C₁₂ and 1,2,3,6,7,8-HxCDD-¹³C₁₂ while 1,2,3,6,7,8-HxCDD-¹³C₁₂ is the RT reference.

Table 5: HRMS Instrumental Descriptor Parameters

| Descriptor | Exact M/Z | M/Z Type | Elemental Composition | Substance | Type | Theoretical | Ion Ratio QC Limits | | |
|------------|-----------|----------|--|--|------------------|-----------------|---------------------|-------|------|
| | | | | | | Ion Ratio | Upper | Lower | |
| 1 | 303.9016 | M | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_4 \text{O}$ | TCDF | Native | 0.77 | 0.65 | 0.89 | |
| | 305.8987 | M+2 | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_3 \text{Cl} \text{O}$ | TCDF | Native | | | | |
| | 315.9419 | M | $^{13}\text{C}_{12} \text{H}_4 \text{Cl}_4 \text{O}$ | TCDF | ^{13}C | 0.77 | 0.65 | 0.89 | |
| | 317.9389 | M+2 | $^{13}\text{C}_{12} \text{H}_4 \text{Cl}_3 \text{Cl} \text{O}$ | TCDF | ^{13}C | | | | |
| | 316.9824 | Lock | $^{12}\text{C}_9 \text{F}_{11}$ | PFK | Lock | | | | |
| | 319.8965 | M | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_4 \text{O}_2$ | TCDD | Native | 0.77 | 0.65 | 0.89 | |
| | 321.8936 | M+2 | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_3 \text{Cl} \text{O}_2$ | TCDD | Native | | | | |
| | 327.8847 | M+8 | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_4 \text{O}_2$ | TCDD | ^{37}Cl | | | | |
| | 331.9368 | M | $^{13}\text{C}_{12} \text{H}_4 \text{Cl}_4 \text{O}_2$ | TCDD | ^{13}C | 0.77 | 0.65 | 0.89 | |
| | 333.9339 | M+2 | $^{13}\text{C}_{12} \text{H}_4 \text{Cl}_3 \text{Cl} \text{O}_2$ | TCDD | ^{13}C | | | | |
| | 339.8597 | M+2 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}$ | PeCDF | Native | 1.55 | 1.32 | 1.78 | |
| | 341.8568 | M+4 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_3 \text{Cl}_2 \text{O}$ | PeCDF | Native | | | | |
| | 351.9 | M+2 | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}$ | PeCDF | ^{13}C | 1.55 | 1.32 | 1.78 | |
| | 353.897 | M+4 | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_3 \text{Cl}_2 \text{O}$ | PeCDF | ^{13}C | | | | |
| | 375.8364 | M+2 | $^{12}\text{C}_{12} \text{H}_4 \text{Cl}_5 \text{Cl} \text{O}$ | HxCDFPE | CI-DPE | | | | |
| | 409.7974 | M+2 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_6 \text{Cl} \text{O}$ | HpCDFPE | CI-DPE | | | | |
| | 2 | 339.8597 | M+2 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}$ | PeCDF | Native | 1.55 | 1.32 | 1.78 |
| | | 341.8568 | M+4 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_3 \text{Cl}_2 \text{O}$ | PeCDF | Native | | | |
| | | 351.9 | M+2 | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}$ | PeCDF | ^{13}C | 1.55 | 1.32 | 1.78 |
| 353.897 | | M+4 | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_3 \text{Cl}_2 \text{O}$ | PeCDF | ^{13}C | | | | |
| 353.8576 | | M | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_5 \text{O}_2$ | PeCDD | Native | 0.63 | 0.54 | 0.72 | |
| 355.8546 | | M+2 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}_2$ | PeCDD | Native | | | | |
| 366.9792 | | Lock | $^{12}\text{C}_{10} \text{F}_{13}$ | PFK | Lock | | | | |
| 365.8978 | | M | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_5 \text{O}_2$ | PeCDD | ^{13}C | 0.63 | 0.54 | 0.72 | |
| 367.8949 | | M+2 | $^{13}\text{C}_{12} \text{H}_3 \text{Cl}_4 \text{Cl} \text{O}_2$ | PeCDD | ^{13}C | | | | |
| 409.7974 | | M+2 | $^{12}\text{C}_{12} \text{H}_3 \text{Cl}_6 \text{Cl} \text{O}$ | HpCDFPE | CI-DPE | | | | |
| 3 | | 373.8207 | M+2 | $^{12}\text{C}_{12} \text{H}_2 \text{Cl}_5 \text{Cl} \text{O}$ | HxCDF | Native | 1.24 | 1.05 | 1.43 |
| | | 375.8178 | M+4 | $^{12}\text{C}_{12} \text{H}_2 \text{Cl}_4 \text{Cl}_2 \text{O}$ | HxCDF | Native | | | |
| | | 380.976 | Lock | $^{12}\text{C}_8 \text{F}_5$ | PFK | Lock | | | |
| | 383.8639 | M | $^{13}\text{C}_{12} \text{H}_2 \text{Cl}_5 \text{O}$ | HxCDF | ^{13}C | 0.51 | 0.43 | 0.59 | |
| | 385.861 | M+2 | $^{13}\text{C}_{12} \text{H}_2 \text{Cl}_5 \text{Cl} \text{O}$ | HxCDF | ^{13}C | | | | |
| | 389.8156 | M+2 | $^{12}\text{C}_{12} \text{H}_2 \text{Cl}_5 \text{Cl} \text{O}_2$ | HxCDD | Native | 1.24 | 1.05 | 1.43 | |
| | 391.8127 | M+4 | $^{12}\text{C}_{12} \text{H}_2 \text{Cl}_4 \text{Cl}_2 \text{O}_2$ | HxCDD | Native | | | | |
| | 401.8559 | M+2 | $^{13}\text{C}_{12} \text{H}_2 \text{Cl}_5 \text{Cl} \text{O}_2$ | HxCDD | ^{13}C | 1.24 | 1.05 | 1.43 | |
| | 403.853 | M+4 | $^{13}\text{C}_{12} \text{H}_2 \text{Cl}_4 \text{Cl}_2 \text{O}_2$ | HxCDD | ^{13}C | | | | |
| | 445.7555 | M+4 | $^{12}\text{C}_{12} \text{H}_2 \text{Cl}_6 \text{Cl}_2 \text{O}$ | OCDFPE | CI-DPE | | | | |
| | 4 | 409.7789 | M+4 | $^{12}\text{C}_{12} \text{H} \text{Cl}_5 \text{Cl}_2 \text{O}$ | HpCDF | Native | 1.88 | 1.60 | 2.16 |
| 411.7759 | | M+6 | $^{12}\text{C}_{12} \text{H} \text{Cl}_4 \text{Cl}_3 \text{O}$ | HpCDF | Native | | | | |
| 417.8253 | | M | $^{13}\text{C}_{12} \text{H} \text{Cl}_7 \text{O}$ | HpCDF | ^{13}C | 0.44 | 0.37 | 0.51 | |
| 419.822 | | M+2 | $^{13}\text{C}_{12} \text{H} \text{Cl}_6 \text{Cl} \text{O}$ | HpCDF | ^{13}C | | | | |
| 423.7767 | | M+2 | $^{12}\text{C}_{12} \text{H} \text{Cl}_6 \text{Cl} \text{O}_2$ | HpCDD | Native | 1.04 | 0.88 | 1.20 | |
| 425.7737 | | M+4 | $^{12}\text{C}_{12} \text{H} \text{Cl}_5 \text{Cl}_2 \text{O}_2$ | HpCDD | Native | | | | |
| 430.9728 | | Lock | $^{12}\text{C}_9 \text{F}_{17}$ | PFK | Lock | | | | |
| 435.8169 | | M+2 | $^{13}\text{C}_{12} \text{H} \text{Cl}_6 \text{Cl} \text{O}_2$ | HpCDD | ^{13}C | 1.04 | 0.88 | 1.20 | |
| 437.814 | | M+4 | $^{13}\text{C}_{12} \text{H} \text{Cl}_5 \text{Cl}_2 \text{O}_2$ | HpCDD | ^{13}C | | | | |
| 479.7165 | | M+4 | $^{12}\text{C}_{12} \text{H} \text{Cl}_7 \text{Cl}_2 \text{O}$ | NCDPE | CI-DPE | | | | |
| 5 | | 441.7428 | M+2 | $^{12}\text{C}_{12} \text{Cl}_7 \text{Cl} \text{O}$ | OCDF | Native | 0.89 | 0.76 | 1.02 |
| | 443.7399 | M+4 | $^{12}\text{C}_{12} \text{Cl}_6 \text{Cl}_2 \text{O}$ | OCDF | Native | | | | |
| | 454.9728 | Lock | $^{12}\text{C}_{11} \text{F}_{17}$ | PFK | Lock | | | | |
| | 457.7377 | M+2 | $^{12}\text{C}_{12} \text{Cl}_7 \text{Cl} \text{O}_2$ | OCDD | Native | 0.89 | 0.76 | 1.02 | |
| | 459.7348 | M+4 | $^{12}\text{C}_{12} \text{Cl}_6 \text{Cl}_2 \text{O}_2$ | OCDD | Native | | | | |
| | 469.778 | M+2 | $^{13}\text{C}_{12} \text{Cl}_7 \text{Cl} \text{O}_2$ | OCDD | ^{13}C | 0.89 | 0.76 | 1.02 | |
| | 471.775 | M+4 | $^{13}\text{C}_{12} \text{Cl}_6 \text{Cl}_2 \text{O}_2$ | OCDD | ^{13}C | | | | |
| | 513.6775 | M+4 | $^{12}\text{C}_{12} \text{Cl}_8 \text{Cl}_2 \text{O}$ | DCDFPE | CI-DPE | | | | |

Data Calculations:

a) Analyte Concentrations:

The relative response factor of each target relative to the standard against which it is to be calculated is determined using the area responses of both quantification ions via equation 9.1.

In cases where a native target is calculated against an exact labelled analogue, the quantification will be considered to be by isotope dilution. In other cases, the quantification will be considered to be by internal standard.

$$\text{RRF} = \frac{(A1_t + A2_t) C_s}{(A1_s + A2_s) C_t} \quad \text{Equ. 9.1}$$

Where,

$A1_t + A2_t$: The areas of the two quantification ions for the target analyte

$A1_s + A2_s$: The areas of the two quantification ions for the labelled compound against which the target analyte will be calculated.

C_t : The concentration in the calibration standard of the target analyte.

C_s : The concentration in the calibration standard of the labelled compound against which the target will be calculated.

For all analytes to be quantified and from the initial calibration series of standard injections, a table of RRFs is prepared. The relative standard deviation (%RSD, or the coefficient of variance) is checked to confirm that the appropriate method criteria has been met as listed in Table 3. The average of the five or six levels of standard for each analyte, RRF_{av} is applied for quantification of samples according to Equations 9.2 and 9.3 below.

$$\text{Amount in sample (pg)} = \frac{(A1_n + A2_n) Q_i}{(A1_i + A2_i) (\text{RRF}_{\text{av}})} \quad \text{Equ. 9.2}$$

$$\text{Concentration in sample (pg/g or pg/l)} = \frac{(A1_n + A2_n) Q_i}{(A1_i + A2_i) (\text{RRF}_{\text{av}}) (W_s)} \quad \text{Equ. 9.3}$$

Where,

Q_i = The amount (pg) of labelled compound added to the sample

W_s = The weight (g) or volume (l) of sample

b) Extraction, Clean-up, and Sampling Standard Recovery Calculation:

The extraction, clean-up, and sampling standard recoveries are determined by Equation 9.4 below.

$$\% \text{ Recovery} = (\text{Amount in sample}) / (\text{Amount added to sample}) \times 100 \quad \text{Equ. 9.4}$$

c) Estimated Detection Limit

$$\text{EDL} = \frac{2.5 \times H_x \times Q_{\text{es}}}{H_{\text{es}} \times W \times \text{RRF}_{\text{av}}} \quad \text{Equ. 9.5}$$

Where,

EDL = estimated detection limit for homologous 2,3,7,8-Substituted PCDD/Fs

H_x = sum of the height of the noise level for each quantification ions for the unlabelled PCDD/Fs.

H_{es} = Sum of the heights of responses of both quantification ions for the labelled extraction standard.

W = weight of volume of sample

RRF_{av} = average relative response factor

Q_{es} = Amount of extraction standard added

Chromatogram Annotation Codes

All manually integrated peaks are expanded and reprinted with the following annotations:

* Analyst Initials AA
 * Date YYMMDD
 * integration code CC

The Syntax is: Example:
 AAYYMMDDCC SK111220MB

| Code | Mnemonic | Description |
|-------|----------------------------|--|
| MB | Manual Baseline | The peak was manually integrated because the initial baseline was determined incorrectly by the software |
| MS | Manual Split | The peak was manually integrated because the peak was incorrectly or not split by the software |
| MJ/MC | Manual Join/Manual Combine | The peak was manually integrated because the peak was split by the software and the peak should be integrated as a single peak |
| MA | Manual Add | The peak was manually integrated because the signal:noise ratio was judged to be >2.5 |
| MD | Manual Delete | The peak was excluded because the signal:noise ratio was judged to be <2.5 |
| MX | Manual Exclude | The peak was excluded due to an interference |
| NH | Noise Height | The noise height for detection limit calculation was manually defined, over-riding the software chosen value |
| MT | Manual Time | The peak retention time was manually chosen |

The following explanatory annotation codes may appear on the chromatograms of peaks that have been reviewed:

| Code | Mnemonic | Description |
|-------|--|---|
| + | Detected Peak | A peak was detected at this mass and retention time that was above 2.5:1 signal to noise |
| < | Below Detection Limit | The signal at this mass and retention time was below 2.5:1 signal to noise |
| EMPC | Estimated Maximum Possible Concentration | The signal at this mass and retention time is an interference such that the target compound could not be confirmed |
| X-RT | Not Detected due to Retention Time non-conformance | The signal at this retention time could not be used to positively identify the target compound because of retention time non-conformance (apex of quantification and confirmation ions do not maximize within the same two seconds, or the retention time of the peak does not fall within the expected range with respect to its labeled analogue) |
| X-LOC | Not Detected due to interference from a higher level of chlorination | The signal at this retention time is attributable to a fragment from a co-eluting compound at a higher level of chlorination, and cannot be used to positively identify the target. The result is expressed as an Estimated Maximum Possible Concentration (EMPC) |
| X-DPE | Not Detected due to diphenyl ether interference | The signal at this retention time is attributable to interference from a chlorinated diphenyl ether, and cannot be used to positively identify the target. The result is expressed as an Estimated Maximum Possible Concentration (EMPC) |
| X-IF | Not Detected due to interference | The signal at this retention time is attributable to a co-eluting interference, and cannot be used to positively identify the target. The result is expressed as an Estimated Maximum Possible Concentration (EMPC) |

Deviations from the Primary Reference Methods:

The following changes and clarifications apply:

application of one standard calibration series to all of these methods is within the scope of each and every one of the methods. The calibration standard set CS1 through CS5 is consistent with the standards concentration listing in method 1613B Table 4. The CS0 extends the calibration range below what is required by all of the methods. Table 4 defines the use of each of the labelled standards relative to each of the methods.

a. Method 1613B lists a larger suite of labelled extraction standards than does method 8290A. Additional labelled extraction standards have been added into the 8290A analysis to enhance the method and the data quality. These additions to the method constitute performance based enhancements and are within the scope of SW846 Method 8290A.

b. The levels presented in the calibration table of method 8290A are recommended values only. Changes to these concentrations, especially to expand the range, are within the scope of the method. Therefore application of the 1613B calibration standards to method 8290A is compliant with the scope of the method.

c. TO-9A is also a performance based method. It specifically states that different extraction standards and different concentrations of standards from those listed in TO-9A Table 3 is acceptable (see Section 6.8 of reference method).

d. Although OAQPS reference method 23 is not a performance based method, application of the 1613B standards has been defined as within the scope of the method. (see Appendix B)

2) Chlorinated Diphenyl Ether interferences: Both methods 1613B and 8290A indicate that any instrumental response showing the presence of a chlorinated diphenyl ether response and that coelutes with a PCDF represents an interference on that analyte (see Sections 18.3 and 7.8.4.4 respectively). This apparent zero tolerance does not take into account that the response in the diphenyl ether channel may be trivial relative to the corresponding PCDF. For this 'Standard Method', we have defined a chlorinated diphenyl ether interference as the presence of a **significant** response within the chlorinated diphenyl ether channel (rather than zero response) and defined significant as a response equal to or greater than 10% of the peak response in the PCDF channel.

3) When the primary analysis is performed using a DB5MS GC column, 2,3,7,8-TCDF can be resolved to a valley height of 60% from the closest-eluting isomers for this column, providing good quantification of this target without further confirmation. Confirmation of 2,3,7,8-TCDF concentrations above the level of the lowest calibration standard are performed on a second column on a contract basis when requested. Confirmation of additional 2,3,7,8-substituted PCDD/F isomers is also available when

4) Although not categorically stated in all associated PCDD/F methods, we maintain that each and every individual clean-up procedure is, by definition, performance-based and optional. There is not an expectation within the industry to follow exactly the descriptions of clean-ups in reference methods. Adaptations which meet or exceed the required performance criteria are therefore acceptable within the scope of each reference method. The reference method descriptions are intended as guidelines or templates available to help the laboratory to define effective in-house clean-up methods. The objective within the laboratory is to provide quality clean extracts to the instrument for analysis. Each individual clean-up is part of the laboratory's 'arsenal' in order to

5) There are differences within the individual reference methods as to the precise spiking protocols for adding extraction standards and native spikes (for LCS, MS and MSD). To ensure consistency within the laboratory between PCDD/F and related methods, the PCDD/F preparative 'Specific Method' requires solids (including stack and ambient sorbants/filters) to be spiked in the soxhlet thimble from a nonane solution and waters are spiked before filtering from an acetone solution. This is consistent with the 8290A

6) Sub-sampling of solids and pre-extraction processing is done in a manner that minimizes potential for cross-contamination. These processes are designed around SW846 protocols rather than 1613B protocols. Solids are sub-sampled directly from the bottle as submitted to the laboratory wherever practical. If the sample is submitted such that homogenization in the bottle is impractical (eg. the bottle is too full or lumps cannot be broken down), then transferring the sample to a tray or another bottle

7) The concentrations of labelled and native spiking solutions are not consistent with those listed in all of the reference methods. These concentrations are prepared at levels convenient and expedient for accurate laboratory processing.

the inter-laboratory performance limits defined in method 1613B rather than the relatively arbitrary limits of 35-140% suggested in Section 8.4 of method 8290A.

9) With respect to ions monitored for P5CDD and H7CDF:

a. The 358 ion has a potential for interference from PCB (hexachlorobiphenyls) dependent upon levels of PCBs in the sample and the instrument tuning. Of particular concern is PCB-169 which on a DB5MS column elutes very close to 1,2,3,7,8-P5CDD and which is not removed for the PCDD/F extracts even by carbon clean-up. To eliminate the potential of such interferences from PCB on the 358 mass, we choose to monitor the alternate ion pair of 354 and 356.

b. Similarly, the 408 ion of native H7CDF is prone to problematic interferences arising from ¹³C¹²-labeled heptachlorinated biphenyls. To eliminate the potential of such interferences from PCB on the 358 mass, we choose to monitor the alternate ion pair

SVOC DATA PACKAGE

SECTION 4: CALIBRATION DATA

Including:

for Multi-Point Calibration(s)

- Multi-Point Calibration Tables
- Individual Quantitation Reports

for Continuing Calibration(s)

- Individual Quantitation Reports

ALS Life Sciences

Calibration Summary Report

| Calibration Level | Filename | Run Date |
|-------------------|-------------|-------------------|
| CS-1 | 7-201001A03 | 01-Oct-2020 11:01 |
| CS-2 | 7-201001A02 | 01-Oct-2020 10:19 |
| CS-3 | 7-201001A01 | 01-Oct-2020 09:38 |
| CS-4 | 7-201001A07 | 01-Oct-2020 13:51 |
| CS-5 | 7-201001A06 | 01-Oct-2020 13:08 |

| | |
|-----------|--|
| Approved: | <i>N Ashtari</i> --e-signature-- 28-Jan-2021 |
|-----------|--|

| Target Analytes | Relative Response Factors | | | | | Mean | % RSD |
|----------------------------------|---------------------------|-------|-------|-------|-------|-------|-------|
| | CS-1 | CS-2 | CS-3 | CS-4 | CS-5 | | |
| 2,3,7,8-TCDD | 0.966 | 0.960 | 1.149 | 1.006 | 0.998 | 1.016 | 8% |
| 1,2,3,7,8-PeCDD | 0.861 | 0.913 | 0.899 | 0.926 | 0.902 | 0.900 | 3% |
| 1,2,3,4,7,8-HxCDD | 0.824 | 0.819 | 0.841 | 0.849 | 0.901 | 0.847 | 4% |
| 1,2,3,6,7,8-HxCDD | 0.988 | 0.942 | 0.959 | 0.967 | 0.947 | 0.961 | 2% |
| 1,2,3,7,8,9-HxCDD | 0.882 | 0.853 | 0.875 | 0.900 | 0.901 | 0.882 | 2% |
| 1,2,3,4,6,7,8-HpCDD | 0.893 | 0.878 | 0.952 | 0.947 | 0.925 | 0.919 | 4% |
| OCDD | 1.008 | 0.968 | 0.923 | 0.994 | 0.924 | 0.963 | 4% |
| 2,3,7,8-TCDF | 0.873 | 0.873 | 0.916 | 0.934 | 0.933 | 0.906 | 3% |
| 1,2,3,7,8-PeCDF | 0.937 | 0.957 | 0.939 | 1.007 | 1.001 | 0.968 | 3% |
| 2,3,4,7,8-PeCDF | 1.001 | 1.019 | 1.042 | 1.071 | 1.063 | 1.039 | 3% |
| 1,2,3,4,7,8-HxCDF | 1.068 | 1.128 | 1.112 | 1.122 | 1.100 | 1.106 | 2% |
| 1,2,3,6,7,8-HxCDF | 1.133 | 1.140 | 1.188 | 1.172 | 1.150 | 1.157 | 2% |
| 2,3,4,6,7,8-HxCDF | 1.003 | 1.051 | 1.053 | 1.081 | 1.070 | 1.052 | 3% |
| 1,2,3,7,8,9-HxCDF | 0.874 | 0.910 | 0.928 | 0.942 | 0.928 | 0.916 | 3% |
| 1,2,3,4,6,7,8-HpCDF | 1.028 | 1.017 | 1.016 | 1.045 | 1.021 | 1.025 | 1% |
| 1,2,3,4,7,8,9-HpCDF | 0.790 | 0.800 | 0.783 | 0.833 | 0.802 | 0.802 | 2% |
| OCDF | 1.246 | 1.299 | 1.233 | 1.351 | 1.215 | 1.269 | 4% |
| Field Spike Standards | | | | | | | |
| 37Cl4-2,3,7,8-TCDD | 0.964 | 1.033 | 1.337 | 0.992 | 1.028 | 1.071 | 14% |
| 13C12-1,2,3,4,7,8-HxCDD | 0.883 | 0.863 | 0.883 | 0.870 | 0.943 | 0.888 | 4% |
| 13C12-2,3,4,7,8-PeCDF | 0.958 | 0.957 | 1.000 | 0.965 | 0.968 | 0.970 | 2% |
| 13C12-1,2,3,4,7,8-HxCDF | 0.894 | 0.922 | 0.923 | 0.898 | 0.891 | 0.906 | 2% |
| 13C12-1,2,3,4,7,8,9-HpCDF | 0.786 | 0.758 | 0.823 | 0.768 | 0.750 | 0.777 | 4% |
| Extraction Standards | | | | | | | |
| 13C12-2,3,7,8-TCDD | 1.096 | 1.110 | 0.972 | 1.120 | 1.237 | 1.107 | 9% |
| 13C12-1,2,3,7,8-PeCDD | 0.719 | 0.730 | 0.787 | 0.769 | 0.918 | 0.785 | 10% |
| 13C12-1,2,3,6,7,8-HxCDD | 0.978 | 1.002 | 1.149 | 1.009 | 0.979 | 1.023 | 7% |
| 13C12-1,2,3,4,6,7,8-HpCDD | 0.690 | 0.700 | 0.736 | 0.695 | 0.713 | 0.707 | 3% |
| 13C12-OCDD | 0.392 | 0.404 | 0.452 | 0.444 | 0.493 | 0.437 | 9% |
| 13C12-2,3,7,8-TCDF | 1.365 | 1.390 | 1.529 | 1.396 | 1.468 | 1.430 | 5% |
| 13C12-1,2,3,7,8-PeCDF | 0.992 | 1.012 | 1.127 | 1.061 | 1.229 | 1.084 | 9% |
| 13C12-1,2,3,6,7,8-HxCDF | 1.195 | 1.203 | 1.402 | 1.199 | 1.189 | 1.238 | 7% |
| 13C12-1,2,3,4,6,7,8-HpCDF | 0.779 | 0.800 | 0.929 | 0.814 | 0.823 | 0.829 | 7% |
| Cleanup Standard | | | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 0.924 | 0.937 | 1.102 | 0.925 | 0.937 | 0.965 | 8% |

ALS Life Sciences

Calibration Report

ALS Sample ID **H7-20-CS1-1209**
 Analysis Method EPA M23
 Analysis Type Calibration

Filename 7-201001A03 Inst # HRMS-7 Column DB5MSUSR188441H Run Date 01-Oct-2020 11:01

Approved: *N Ashtari*
 --e-signature--
 28-Jan-2021

| Target Analytes | Ret. Time | Ion Ratio | Concentration ng/mL | Response | RRF |
|----------------------------------|-----------|-----------|---------------------|-----------|-----------|
| 2,3,7,8-TCDD | 27.85 | 0.81 | 0.50 | 1.02E+04 | 0.966 |
| 1,2,3,7,8-PeCDD | 32.29 | 1.66 | 2.50 | 2.99E+04 | 0.861 |
| 1,2,3,4,7,8-HxCDD | 34.38 | 1.22 | 2.50 | 2.71E+04 | 0.824 |
| 1,2,3,6,7,8-HxCDD | 34.44 | 1.27 | 2.50 | 3.25E+04 | 0.988 |
| 1,2,3,7,8,9-HxCDD | 34.56 | 1.22 | 2.50 | 2.90E+04 | 0.882 |
| 1,2,3,4,6,7,8-HpCDD | 36 | 1.06 | 2.50 | 2.07E+04 | 0.893 |
| OCDD | 37.43 | 0.90 | 5.00 | 2.65E+04 | 1.008 |
| 2,3,7,8-TCDF | 26.95 | 0.80 | 0.50 | 1.15E+04 | 0.873 |
| 1,2,3,7,8-PeCDF | 31.29 | 1.54 | 2.50 | 4.49E+04 | 0.937 |
| 2,3,4,7,8-PeCDF | 32.07 | 1.50 | 2.50 | 4.80E+04 | 1.001 |
| 1,2,3,4,7,8-HxCDF | 33.88 | 1.26 | 2.50 | 4.28E+04 | 1.068 |
| 1,2,3,6,7,8-HxCDF | 33.95 | 1.22 | 2.50 | 4.55E+04 | 1.133 |
| 2,3,4,6,7,8-HxCDF | 34.29 | 1.25 | 2.50 | 4.02E+04 | 1.003 |
| 1,2,3,7,8,9-HxCDF | 34.71 | 1.30 | 2.50 | 3.50E+04 | 0.874 |
| 1,2,3,4,6,7,8-HpCDF | 35.47 | 2.03 | 2.50 | 2.69E+04 | 1.028 |
| 1,2,3,4,7,8,9-HpCDF | 36.25 | 1.90 | 2.50 | 2.06E+04 | 0.790 |
| OCDF | 37.52 | 0.92 | 5.00 | 3.28E+04 | 1.246 |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 27.85 | 0.00 | 0.50 | 1.02E+04 | 0.964 |
| 13C12-1,2,3,4,7,8-HxCDD | 34.38 | 1.28 | 100.00 | 1.16E+06 | 0.883 |
| 13C12-2,3,4,7,8-PeCDF | 32.05 | 1.51 | 100.00 | 1.84E+06 | 0.958 |
| 13C12-1,2,3,4,7,8-HxCDF | 33.87 | 0.53 | 100.00 | 1.44E+06 | 0.894 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 36.25 | 0.44 | 100.00 | 8.22E+05 | 0.786 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 27.82 | 0.784 | 100 | 2.12E+06 | 1.096 |
| 13C12-1,2,3,7,8-PeCDD | 32.28 | 1.602 | 100 | 1.39E+06 | 0.719 |
| 13C12-1,2,3,6,7,8-HxCDD | 34.43 | 1.3 | 100 | 1.31E+06 | 0.978 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 36 | 1.056 | 100 | 9.27E+05 | 0.69 |
| 13C12-OCDD | 37.42 | 0.883 | 200 | 1.05E+06 | 0.392 |
| 13C12-2,3,7,8-TCDF | 26.93 | 0.757 | 100 | 2.64E+06 | 1.365 |
| 13C12-1,2,3,7,8-PeCDF | 31.28 | 1.563 | 100 | 1.92E+06 | 0.992 |
| 13C12-1,2,3,6,7,8-HxCDF | 33.94 | 0.52 | 100 | 1.60E+06 | 1.195 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 35.46 | 0.464 | 100 | 1.05E+06 | 0.779 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 34.7 | 0.522 | 100 | 1.24E+06 | 0.924 |
| Injection Standards | | | | | |
| 13C12-1234-TCDD IS | 27.13 | 0.792 | 100 | 1933155.7 | 19331.557 |
| 13C12-123789-HxCDD IS | 34.56 | 1.26 | 100.00 | 1.34E+06 | 13434.236 |

ALS Life Sciences

Calibration Report

ALS Sample ID **H7-20-CS2-1209**
 Analysis Method EPA M23
 Analysis Type Calibration

Filename: 7-201001A02 Inst #: HRMS-7 Column: DB5MSUSR188441H Run Date: 01-Oct-2020 10:19

Approved: *N Ashtari*
 --e-signature--
 28-Jan-2021

| Target Analytes | Ret. Time | Ion Ratio | Concentration ng/mL | Response | RRF |
|----------------------------------|-----------|-----------|---------------------|-----------|-----------|
| 2,3,7,8-TCDD | 27.83 | 0.79 | 2.00 | 3.80E+04 | 0.960 |
| 1,2,3,7,8-PeCDD | 32.28 | 1.61 | 10.00 | 1.19E+05 | 0.913 |
| 1,2,3,4,7,8-HxCDD | 34.37 | 1.27 | 10.00 | 1.01E+05 | 0.819 |
| 1,2,3,6,7,8-HxCDD | 34.43 | 1.33 | 10.00 | 1.17E+05 | 0.942 |
| 1,2,3,7,8,9-HxCDD | 34.55 | 1.30 | 10.00 | 1.06E+05 | 0.853 |
| 1,2,3,4,6,7,8-HpCDD | 35.99 | 1.02 | 10.00 | 7.60E+04 | 0.878 |
| OCDD | 37.42 | 0.87 | 20.00 | 9.66E+04 | 0.968 |
| 2,3,7,8-TCDF | 26.95 | 0.80 | 2.00 | 4.33E+04 | 0.873 |
| 1,2,3,7,8-PeCDF | 31.28 | 1.62 | 10.00 | 1.73E+05 | 0.957 |
| 2,3,4,7,8-PeCDF | 32.06 | 1.58 | 10.00 | 1.84E+05 | 1.019 |
| 1,2,3,4,7,8-HxCDF | 33.86 | 1.17 | 10.00 | 1.68E+05 | 1.128 |
| 1,2,3,6,7,8-HxCDF | 33.94 | 1.14 | 10.00 | 1.70E+05 | 1.140 |
| 2,3,4,6,7,8-HxCDF | 34.28 | 1.18 | 10.00 | 1.56E+05 | 1.051 |
| 1,2,3,7,8,9-HxCDF | 34.7 | 1.18 | 10.00 | 1.35E+05 | 0.910 |
| 1,2,3,4,6,7,8-HpCDF | 35.45 | 2.01 | 10.00 | 1.01E+05 | 1.017 |
| 1,2,3,4,7,8,9-HpCDF | 36.23 | 1.99 | 10.00 | 7.91E+04 | 0.800 |
| OCDF | 37.5 | 0.90 | 20.00 | 1.30E+05 | 1.299 |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 27.83 | 0.00 | 2.00 | 4.09E+04 | 1.033 |
| 13C12-1,2,3,4,7,8-HxCDD | 34.36 | 1.24 | 100.00 | 1.07E+06 | 0.863 |
| 13C12-2,3,4,7,8-PeCDF | 32.05 | 1.57 | 100.00 | 1.73E+06 | 0.957 |
| 13C12-1,2,3,4,7,8-HxCDF | 33.86 | 0.53 | 100.00 | 1.37E+06 | 0.922 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 36.23 | 0.47 | 100.00 | 7.49E+05 | 0.758 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 27.81 | 0.781 | 100 | 1.98E+06 | 1.11 |
| 13C12-1,2,3,7,8-PeCDD | 32.26 | 1.612 | 100 | 1.30E+06 | 0.73 |
| 13C12-1,2,3,6,7,8-HxCDD | 34.42 | 1.245 | 100 | 1.24E+06 | 1.002 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 35.99 | 1.077 | 100 | 8.66E+05 | 0.7 |
| 13C12-OCDD | 37.41 | 0.925 | 200 | 9.99E+05 | 0.404 |
| 13C12-2,3,7,8-TCDF | 26.92 | 0.769 | 100 | 2.48E+06 | 1.39 |
| 13C12-1,2,3,7,8-PeCDF | 31.27 | 1.564 | 100 | 1.80E+06 | 1.012 |
| 13C12-1,2,3,6,7,8-HxCDF | 33.93 | 0.518 | 100 | 1.49E+06 | 1.203 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 35.45 | 0.452 | 100 | 9.89E+05 | 0.8 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 34.69 | 0.517 | 100 | 1.16E+06 | 0.937 |
| Injection Standards | | | | | |
| 13C12-1234-TCDD IS | 27.11 | 0.788 | 100 | 1784134.8 | 17841.348 |
| 13C12-123789-HxCDD IS | 34.54 | 1.24 | 100.00 | 1.24E+06 | 12368.686 |

ALS Life Sciences

Calibration Report

ALS Sample ID **H7-20-CCV-1209**
 Analysis Method EPA M23
 Analysis Type Calibration

Filename: 7-201001A01 Inst #: HRMS-7 Column: DB5MSUSR188441H Run Date: 01-Oct-2020 09:38

Approved: *N Ashtari*
 --e-signature--
 28-Jan-2021

| Target Analytes | Ret. Time | Ion Ratio | Concentration ng/mL | Response | RRF |
|----------------------------------|-----------|-----------|---------------------|-----------|-----------|
| 2,3,7,8-TCDD | 27.87 | 0.77 | 10.00 | 1.68E+05 | 1.149 |
| 1,2,3,7,8-PeCDD | 32.29 | 1.63 | 50.00 | 5.31E+05 | 0.899 |
| 1,2,3,4,7,8-HxCDD | 34.38 | 1.23 | 50.00 | 4.59E+05 | 0.841 |
| 1,2,3,6,7,8-HxCDD | 34.43 | 1.21 | 50.00 | 5.24E+05 | 0.959 |
| 1,2,3,7,8,9-HxCDD | 34.55 | 1.21 | 50.00 | 4.77E+05 | 0.875 |
| 1,2,3,4,6,7,8-HpCDD | 36 | 1.06 | 50.00 | 3.33E+05 | 0.952 |
| OCDD | 37.42 | 0.93 | 100.00 | 3.97E+05 | 0.923 |
| 2,3,7,8-TCDF | 26.98 | 0.80 | 10.00 | 2.10E+05 | 0.916 |
| 1,2,3,7,8-PeCDF | 31.29 | 1.59 | 50.00 | 7.95E+05 | 0.939 |
| 2,3,4,7,8-PeCDF | 32.07 | 1.58 | 50.00 | 8.82E+05 | 1.042 |
| 1,2,3,4,7,8-HxCDF | 33.88 | 1.21 | 50.00 | 7.41E+05 | 1.112 |
| 1,2,3,6,7,8-HxCDF | 33.95 | 1.22 | 50.00 | 7.91E+05 | 1.188 |
| 2,3,4,6,7,8-HxCDF | 34.28 | 1.20 | 50.00 | 7.02E+05 | 1.053 |
| 1,2,3,7,8,9-HxCDF | 34.7 | 1.21 | 50.00 | 6.19E+05 | 0.928 |
| 1,2,3,4,6,7,8-HpCDF | 35.46 | 1.97 | 50.00 | 4.49E+05 | 1.016 |
| 1,2,3,4,7,8,9-HpCDF | 36.24 | 1.88 | 50.00 | 3.46E+05 | 0.783 |
| OCDF | 37.51 | 0.94 | 100.00 | 5.30E+05 | 1.233 |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 27.87 | 0.00 | 10.00 | 1.95E+05 | 1.337 |
| 13C12-1,2,3,4,7,8-HxCDD | 34.37 | 1.25 | 100.00 | 9.64E+05 | 0.883 |
| 13C12-2,3,4,7,8-PeCDF | 32.06 | 1.54 | 100.00 | 1.69E+06 | 1.000 |
| 13C12-1,2,3,4,7,8-HxCDF | 33.87 | 0.52 | 100.00 | 1.23E+06 | 0.923 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 36.24 | 0.46 | 100.00 | 7.27E+05 | 0.823 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 27.84 | 0.799 | 100 | 1.46E+06 | 0.972 |
| 13C12-1,2,3,7,8-PeCDD | 32.28 | 1.607 | 100 | 1.18E+06 | 0.787 |
| 13C12-1,2,3,6,7,8-HxCDD | 34.43 | 1.257 | 100 | 1.09E+06 | 1.149 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 36 | 1.028 | 100 | 7.00E+05 | 0.736 |
| 13C12-OCDD | 37.42 | 0.861 | 200 | 8.60E+05 | 0.452 |
| 13C12-2,3,7,8-TCDF | 26.95 | 0.771 | 100 | 2.30E+06 | 1.529 |
| 13C12-1,2,3,7,8-PeCDF | 31.28 | 1.551 | 100 | 1.69E+06 | 1.127 |
| 13C12-1,2,3,6,7,8-HxCDF | 33.94 | 0.521 | 100 | 1.33E+06 | 1.402 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 35.46 | 0.451 | 100 | 8.83E+05 | 0.929 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 34.7 | 0.533 | 100 | 1.05E+06 | 1.102 |
| Injection Standards | | | | | |
| 13C12-1234-TCDD IS | 27.16 | 0.786 | 100 | 1502322.7 | 15023.227 |
| 13C12-123789-HxCDD IS | 34.55 | 1.31 | 100.00 | 9.51E+05 | 9505.725 |

ALS Life Sciences

Calibration Report

ALS Sample ID **H7-20-CS4-1209**
 Analysis Method EPA M23
 Analysis Type Calibration

Filename: 7-201001A07 Inst #: HRMS-7 Column: DB5MSUSR188441H Run Date: 01-Oct-2020 13:51

Approved: *N Ashtari*
 --e-signature--
 28-Jan-2021

| Target Analytes | Ret. Time | Ion Ratio | Concentration ng/mL | Response | RRF |
|----------------------------------|-----------|-----------|---------------------|-----------|-----------|
| 2,3,7,8-TCDD | 27.84 | 0.78 | 40.00 | 7.02E+05 | 1.006 |
| 1,2,3,7,8-PeCDD | 32.28 | 1.63 | 200.00 | 2.22E+06 | 0.926 |
| 1,2,3,4,7,8-HxCDD | 34.37 | 1.22 | 200.00 | 2.01E+06 | 0.849 |
| 1,2,3,6,7,8-HxCDD | 34.43 | 1.23 | 200.00 | 2.29E+06 | 0.967 |
| 1,2,3,7,8,9-HxCDD | 34.55 | 1.23 | 200.00 | 2.13E+06 | 0.900 |
| 1,2,3,4,6,7,8-HpCDD | 36 | 1.05 | 200.00 | 1.55E+06 | 0.947 |
| OCDD | 37.42 | 0.89 | 400.00 | 2.07E+06 | 0.994 |
| 2,3,7,8-TCDF | 26.95 | 0.78 | 40.00 | 8.12E+05 | 0.934 |
| 1,2,3,7,8-PeCDF | 31.28 | 1.59 | 200.00 | 3.33E+06 | 1.007 |
| 2,3,4,7,8-PeCDF | 32.06 | 1.57 | 200.00 | 3.54E+06 | 1.071 |
| 1,2,3,4,7,8-HxCDF | 33.87 | 1.21 | 200.00 | 3.16E+06 | 1.122 |
| 1,2,3,6,7,8-HxCDF | 33.94 | 1.21 | 200.00 | 3.30E+06 | 1.172 |
| 2,3,4,6,7,8-HxCDF | 34.28 | 1.19 | 200.00 | 3.04E+06 | 1.081 |
| 1,2,3,7,8,9-HxCDF | 34.7 | 1.22 | 200.00 | 2.65E+06 | 0.942 |
| 1,2,3,4,6,7,8-HpCDF | 35.46 | 1.87 | 200.00 | 2.00E+06 | 1.045 |
| 1,2,3,4,7,8,9-HpCDF | 36.24 | 1.90 | 200.00 | 1.59E+06 | 0.833 |
| OCDF | 37.51 | 0.91 | 400.00 | 2.82E+06 | 1.351 |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 27.84 | 0.00 | 40.00 | 6.92E+05 | 0.992 |
| 13C12-1,2,3,4,7,8-HxCDD | 34.37 | 1.25 | 100.00 | 1.03E+06 | 0.870 |
| 13C12-2,3,4,7,8-PeCDF | 32.05 | 1.54 | 100.00 | 1.60E+06 | 0.965 |
| 13C12-1,2,3,4,7,8-HxCDF | 33.86 | 0.53 | 100.00 | 1.26E+06 | 0.898 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 36.23 | 0.44 | 100.00 | 7.35E+05 | 0.768 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 27.81 | 0.77 | 100 | 1.74E+06 | 1.12 |
| 13C12-1,2,3,7,8-PeCDD | 32.27 | 1.59 | 100 | 1.20E+06 | 0.769 |
| 13C12-1,2,3,6,7,8-HxCDD | 34.43 | 1.23 | 100 | 1.18E+06 | 1.009 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 35.99 | 1.046 | 100 | 8.16E+05 | 0.695 |
| 13C12-OCDD | 37.41 | 0.86 | 200 | 1.04E+06 | 0.444 |
| 13C12-2,3,7,8-TCDF | 26.92 | 0.771 | 100 | 2.17E+06 | 1.396 |
| 13C12-1,2,3,7,8-PeCDF | 31.27 | 1.564 | 100 | 1.65E+06 | 1.061 |
| 13C12-1,2,3,6,7,8-HxCDF | 33.93 | 0.524 | 100 | 1.41E+06 | 1.199 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 35.44 | 0.444 | 100 | 9.56E+05 | 0.814 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 34.69 | 0.534 | 100 | 1.09E+06 | 0.925 |
| Injection Standards | | | | | |
| 13C12-1234-TCDD IS | 27.13 | 0.772 | 100 | 1557536.4 | 15575.364 |
| 13C12-123789-HxCDD IS | 34.55 | 1.23 | 100.00 | 1.17E+06 | 11743.281 |

ALS Life Sciences

Calibration Report

ALS Sample ID **H7-20-CS5-1209**
 Analysis Method EPA M23
 Analysis Type Calibration

Filename 7-201001A06 Inst # HRMS-7 Column DB5MSUSR188441H Run Date 01-Oct-2020 13:08

Approved: *N Ashtari*
 --e-signature--
 28-Jan-2021

| Target Analytes | Ret. Time | Ion Ratio | Concentration ng/mL | Response | RRF |
|----------------------------------|-----------|-----------|---------------------|-----------|-----------|
| 2,3,7,8-TCDD | 27.84 | 0.78 | 200.00 | 4.45E+06 | 0.998 |
| 1,2,3,7,8-PeCDD | 32.28 | 1.63 | 1000.00 | 1.49E+07 | 0.902 |
| 1,2,3,4,7,8-HxCDD | 34.37 | 1.23 | 1000.00 | 1.42E+07 | 0.901 |
| 1,2,3,6,7,8-HxCDD | 34.43 | 1.23 | 1000.00 | 1.49E+07 | 0.947 |
| 1,2,3,7,8,9-HxCDD | 34.55 | 1.24 | 1000.00 | 1.42E+07 | 0.901 |
| 1,2,3,4,6,7,8-HpCDD | 36 | 1.06 | 1000.00 | 1.06E+07 | 0.925 |
| OCDD | 37.43 | 0.89 | 2000.00 | 1.47E+07 | 0.924 |
| 2,3,7,8-TCDF | 26.93 | 0.78 | 200.00 | 4.94E+06 | 0.933 |
| 1,2,3,7,8-PeCDF | 31.29 | 1.58 | 1000.00 | 2.22E+07 | 1.001 |
| 2,3,4,7,8-PeCDF | 32.06 | 1.57 | 1000.00 | 2.35E+07 | 1.063 |
| 1,2,3,4,7,8-HxCDF | 33.88 | 1.21 | 1000.00 | 2.11E+07 | 1.100 |
| 1,2,3,6,7,8-HxCDF | 33.95 | 1.22 | 1000.00 | 2.20E+07 | 1.150 |
| 2,3,4,6,7,8-HxCDF | 34.28 | 1.20 | 1000.00 | 2.05E+07 | 1.070 |
| 1,2,3,7,8,9-HxCDF | 34.7 | 1.22 | 1000.00 | 1.78E+07 | 0.928 |
| 1,2,3,4,6,7,8-HpCDF | 35.46 | 1.89 | 1000.00 | 1.35E+07 | 1.021 |
| 1,2,3,4,7,8,9-HpCDF | 36.24 | 1.90 | 1000.00 | 1.06E+07 | 0.802 |
| OCDF | 37.51 | 0.92 | 2000.00 | 1.93E+07 | 1.215 |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 27.83 | 0.00 | 200.00 | 4.58E+06 | 1.028 |
| 13C12-1,2,3,4,7,8-HxCDD | 34.37 | 1.25 | 100.00 | 1.49E+06 | 0.943 |
| 13C12-2,3,4,7,8-PeCDF | 32.05 | 1.54 | 100.00 | 2.14E+06 | 0.968 |
| 13C12-1,2,3,4,7,8-HxCDF | 33.86 | 0.52 | 100.00 | 1.71E+06 | 0.891 |
| 13C12-1,2,3,4,7,8,9-HpCDF | 36.23 | 0.46 | 100.00 | 9.93E+05 | 0.750 |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 27.81 | 0.788 | 100 | 2.23E+06 | 1.237 |
| 13C12-1,2,3,7,8-PeCDD | 32.27 | 1.617 | 100 | 1.65E+06 | 0.918 |
| 13C12-1,2,3,6,7,8-HxCDD | 34.43 | 1.254 | 100 | 1.58E+06 | 0.979 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 35.99 | 1.048 | 100 | 1.15E+06 | 0.713 |
| 13C12-OCDD | 37.42 | 0.886 | 200 | 1.59E+06 | 0.493 |
| 13C12-2,3,7,8-TCDF | 26.92 | 0.772 | 100 | 2.64E+06 | 1.468 |
| 13C12-1,2,3,7,8-PeCDF | 31.28 | 1.553 | 100 | 2.21E+06 | 1.229 |
| 13C12-1,2,3,6,7,8-HxCDF | 33.93 | 0.533 | 100 | 1.92E+06 | 1.189 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 35.46 | 0.458 | 100 | 1.32E+06 | 0.823 |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 34.7 | 0.538 | 100 | 1.51E+06 | 0.937 |
| Injection Standards | | | | | |
| 13C12-1234-TCDD IS | 27.13 | 0.791 | 100 | 1801426.6 | 18014.266 |
| 13C12-123789-HxCDD IS | 34.55 | 1.23 | 100.00 | 1.61E+06 | 16108.986 |

ALS Life Sciences

Second Source Calibration Verification Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CVS | Sampling Date | n/a |
| ALS Sample ID | H7-20-RS1-1209 | Extraction Date | n/a |
| Analysis Method | EPA M23 | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

| |
|---|
| Approved: <i>N Ashtari</i> ---e-signature--- 25-Jan-2021 |
|---|

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-201001A08 |
| Run Date | 01-Oct-20 14:33 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUSR188441H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.84 | 90 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.29 | 106 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.39 | 101 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.44 | 94 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.56 | 102 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 36.01 | 98 | 75-125 | |
| OCDD | 100 | 37.43 | 100 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.95 | 100 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.29 | 106 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 32.07 | 98 | 75-125 | M |
| 1,2,3,4,7,8-HxCDF | 50 | 33.88 | 100 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.95 | 101 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.29 | 100 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.71 | 109 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.47 | 101 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.25 | 108 | 75-125 | |
| OCDF | 100 | 37.52 | 102 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.84 | 107 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.37 | 104 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 32.06 | 100 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.87 | 102 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.24 | 102 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.81 | 101 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.28 | 94 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.43 | 94 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 36.00 | 102 | 70-130 | |
| 13C12-OCDD | 200 | 37.42 | 105 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.92 | 97 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.28 | 93 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.94 | 93 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.46 | 95 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.70 | 95 | 40-130 | |

M Indicates that a peak has been manually integrated.

ALS Life Sciences

Continuing Calibration Report

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|--------------------|----------------|------------------|-------|--|
| Sample Name | CCV | Sampling Date | n/a | Approved: <i>N Ashtari</i> ---e-signature-- 28-Jan-2021 |
| ALS Sample ID | H7-21-CCV-0030 | Extraction Date | n/a | |
| Analysis Method | T09A | Sample Size | 1 n/a | |
| Analysis Type | CCV | Percent Moisture | n/a | |
| Sample Matrix | QC | Split Ratio | 1 | |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210117A32 |
| Run Date | 18-Jan-21 05:21 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.69 | 94 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.08 | 103 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.17 | 100 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.23 | 97 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.35 | 102 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.81 | 100 | 75-125 | |
| OCDD | 100 | 37.27 | 106 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.77 | 100 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.10 | 89 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.86 | 90 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.67 | 89 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.74 | 90 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.08 | 90 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.50 | 93 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.26 | 93 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.06 | 99 | 75-125 | |
| OCDF | 100 | 37.36 | 85 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.69 | 90 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.17 | 103 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.86 | 101 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.66 | 101 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.05 | 99 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.66 | 104 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.07 | 95 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.22 | 96 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.81 | 104 | 70-130 | |
| 13C12-OCDD | 200 | 37.26 | 127 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.75 | 97 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.09 | 95 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.73 | 92 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.25 | 94 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.49 | 96 | 40-130 | |

ALS Life Sciences

Continuing Calibration Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CCV | Sampling Date | n/a |
| ALS Sample ID | H7-21-CCV-0029 | Extraction Date | n/a |
| Analysis Method | T09A | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

Approved:
N Ashtari
 ---e-signature---
 28-Jan-2021

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210117A15 |
| Run Date | 17-Jan-21 17:18 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.69 | 102 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.09 | 105 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.18 | 110 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.23 | 99 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.35 | 106 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.82 | 104 | 75-125 | |
| OCDD | 100 | 37.27 | 109 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.78 | 102 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.12 | 91 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.87 | 91 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.68 | 94 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.75 | 91 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.08 | 94 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.50 | 92 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.26 | 98 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.06 | 99 | 75-125 | |
| OCDF | 100 | 37.36 | 88 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.71 | 91 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.17 | 105 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.86 | 99 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.67 | 102 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.06 | 103 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.68 | 105 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.08 | 99 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.23 | 96 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.81 | 107 | 70-130 | |
| 13C12-OCDD | 200 | 37.27 | 120 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.77 | 105 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.10 | 105 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.74 | 99 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.26 | 101 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.49 | 101 | 40-130 | |

ALS Life Sciences

Continuing Calibration Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CCV | Sampling Date | n/a |
| ALS Sample ID | H7-21-CCV-0041 | Extraction Date | n/a |
| Analysis Method | T09A | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

| |
|---|
| Approved: <i>N Ashtari</i> ---e-signature--- 28-Jan-2021 |
|---|

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210122A01 |
| Run Date | 22-Jan-21 09:56 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.67 | 101 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.08 | 109 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.16 | 103 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.22 | 118 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.35 | 120 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.81 | 111 | 75-125 | |
| OCDD | 100 | 37.26 | 116 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.77 | 105 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.10 | 98 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.85 | 94 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.67 | 96 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.74 | 101 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.07 | 101 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.49 | 95 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.25 | 107 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.05 | 103 | 75-125 | |
| OCDF | 100 | 37.35 | 101 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.67 | 91 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.15 | 105 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.84 | 98 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.66 | 95 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.04 | 96 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.66 | 104 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.06 | 94 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.21 | 95 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.80 | 99 | 70-130 | |
| 13C12-OCDD | 200 | 37.26 | 109 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.74 | 108 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.09 | 106 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.73 | 108 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.25 | 102 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.48 | 104 | 40-130 | |

ALS Life Sciences

Continuing Calibration Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CCV | Sampling Date | n/a |
| ALS Sample ID | H7-21-CCV-0042 | Extraction Date | n/a |
| Analysis Method | TO9A | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

| |
|---|
| Approved: <i>N Ashtari</i> ---e-signature--- 28-Jan-2021 |
|---|

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210122A16 |
| Run Date | 22-Jan-21 20:25 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.67 | 103 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.08 | 110 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.18 | 93 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.23 | 104 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.36 | 103 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.82 | 112 | 75-125 | |
| OCDD | 100 | 37.28 | 119 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.75 | 108 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.10 | 97 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.86 | 96 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.68 | 96 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.74 | 102 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.08 | 102 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.50 | 90 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.26 | 108 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.06 | 102 | 75-125 | |
| OCDF | 100 | 37.36 | 104 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.67 | 92 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.16 | 82 | 75-125 | M |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.85 | 99 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.67 | 96 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.06 | 93 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.64 | 104 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.07 | 89 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.22 | 107 | 75-125 | M |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.81 | 94 | 70-130 | |
| 13C12-OCDD | 200 | 37.27 | 101 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.74 | 109 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.09 | 103 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.73 | 111 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.26 | 99 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.49 | 99 | 40-130 | |

M Indicates that a peak has been manually integrated.

ALS Life Sciences

Continuing Calibration Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CCV | Sampling Date | n/a |
| ALS Sample ID | H7-21-CCV-0457 | Extraction Date | n/a |
| Analysis Method | T09A | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

Approved:
N Ashtari
 ---e-signature---
 28-Jan-2021

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A11 |
| Run Date | 28-Jan-21 00:04 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|--------------|-----------|--------------|---------------|-------|
| 2,3,7,8-TCDD | 10 | 27.66 | 100 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.07 | 104 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.16 | 103 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.22 | 109 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.33 | 116 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.80 | 107 | 75-125 | |
| OCDD | 100 | 37.24 | 108 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.75 | 118 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.09 | 95 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.85 | 96 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.67 | 101 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.73 | 103 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.07 | 107 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.48 | 106 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.25 | 104 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.04 | 111 | 75-125 | |
| OCDF | 100 | 37.33 | 100 | 70-130 | |
| Field Spike Standards | pg/uL | | % Rec | Limits | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.66 | 91 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.15 | 93 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.84 | 98 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.66 | 100 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.04 | 108 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.63 | 103 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.06 | 94 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.21 | 99 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.79 | 115 | 70-130 | |
| 13C12-OCDD | 200 | 37.24 | 165 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.72 | 101 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.07 | 101 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.72 | 103 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.24 | 113 | 70-130 | |
| Cleanup Standard | pg/uL | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.47 | 108 | 40-130 | |

ALS Life Sciences

Continuing Calibration Report

| | | | |
|--------------------|----------------|------------------|-------|
| Sample Name | CCV | Sampling Date | n/a |
| ALS Sample ID | H7-21-CCV-0456 | Extraction Date | n/a |
| Analysis Method | T09A | Sample Size | 1 n/a |
| Analysis Type | CCV | Percent Moisture | n/a |
| Sample Matrix | QC | Split Ratio | 1 |

| |
|---|
| Approved: <i>N Ashtari</i> ---e-signature--- 28-Jan-2021 |
|---|

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A01 |
| Run Date | 27-Jan-21 17:05 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg/uL | Ret. Time | % Rec | Limits | Flags |
|------------------------------|-------|-----------|-------|--------|-------|
| 2,3,7,8-TCDD | 10 | 27.66 | 98 | 75-125 | |
| 1,2,3,7,8-PeCDD | 50 | 32.06 | 106 | 75-125 | |
| 1,2,3,4,7,8-HxCDD | 50 | 34.15 | 105 | 75-125 | |
| 1,2,3,6,7,8-HxCDD | 50 | 34.21 | 113 | 75-125 | |
| 1,2,3,7,8,9-HxCDD | 50 | 34.33 | 112 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDD | 50 | 35.79 | 105 | 75-125 | |
| OCDD | 100 | 37.24 | 110 | 75-125 | |
| 2,3,7,8-TCDF | 10 | 26.75 | 106 | 75-125 | |
| 1,2,3,7,8-PeCDF | 50 | 31.09 | 99 | 75-125 | |
| 2,3,4,7,8-PeCDF | 50 | 31.84 | 96 | 75-125 | |
| 1,2,3,4,7,8-HxCDF | 50 | 33.66 | 102 | 75-125 | |
| 1,2,3,6,7,8-HxCDF | 50 | 33.72 | 105 | 75-125 | |
| 2,3,4,6,7,8-HxCDF | 50 | 34.06 | 107 | 75-125 | |
| 1,2,3,7,8,9-HxCDF | 50 | 34.48 | 107 | 75-125 | |
| 1,2,3,4,6,7,8-HpCDF | 50 | 35.24 | 103 | 75-125 | |
| 1,2,3,4,7,8,9-HpCDF | 50 | 36.03 | 111 | 75-125 | |
| OCDF | 100 | 37.33 | 102 | 70-130 | |
| Field Spike Standards | | | | | |
| 37Cl4-2,3,7,8-TCDD | 10 | 27.66 | 90 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDD | 100 | 34.14 | 101 | 75-125 | |
| 13C12-2,3,4,7,8-PeCDF | 100 | 31.83 | 99 | 75-125 | |
| 13C12-1,2,3,4,7,8-HxCDF | 100 | 33.65 | 99 | 75-125 | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 100 | 36.03 | 105 | 75-125 | |
| Extraction Standards | | | | | |
| 13C12-2,3,7,8-TCDD | 100 | 27.63 | 106 | 75-125 | |
| 13C12-1,2,3,7,8-PeCDD | 100 | 32.05 | 97 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDD | 100 | 34.20 | 98 | 75-125 | |
| 13C12-1,2,3,4,6,7,8-HpCDD | 100 | 35.79 | 116 | 70-130 | |
| 13C12-OCDD | 200 | 37.23 | 150 | 70-130 | |
| 13C12-2,3,7,8-TCDF | 100 | 26.72 | 105 | 70-130 | |
| 13C12-1,2,3,7,8-PeCDF | 100 | 31.07 | 104 | 70-130 | |
| 13C12-1,2,3,6,7,8-HxCDF | 100 | 33.72 | 106 | 70-130 | |
| 13C12-1,2,3,4,6,7,8-HpCDF | 100 | 35.23 | 115 | 70-130 | |
| Cleanup Standard | | | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 100 | 34.47 | 111 | 40-130 | |

SVOC DATA PACKAGE

SECTION 5: QC SAMPLE DATA

Including:

- Laboratory Method Blank Analysis Reports
- Laboratory Control Sample Analysis Reports
- Matrix Spike Analysis Reports
- Other QC Sample Analysis Reports (where applicable)

ALS Life Sciences

Laboratory Method Blank Analysis Report

| | | | | | |
|--------------------|--------------|------------------|-----------|--------|------------------------|
| Sample Name | Method Blank | Sampling Date | n/a | | |
| ALS Sample ID | WG3463499-1 | Extraction Date | 23-Dec-21 | | |
| Analysis Method | T09A | Sample Size | 1 | Sample | Approved: N Ashtari |
| Analysis Type | Blank | Percent Moisture | n/a | | --e-signature-- |
| Sample Matrix | MEDIA | Split Ratio | 4 | | 28-Jan-2021 |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A07 |
| Run Date | 27-Jan-21 21:16 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------------|------------|-----|
| 2,3,7,8-TCDD | 1 | NotFnd | <2.9 | 2.9 | U | 20 |
| 1,2,3,7,8-PeCDD | 1 | NotFnd | <1.7 | 1.7 | U | 100 |
| 1,2,3,4,7,8-HxCDD | 0.1 | NotFnd | <1.0 | 1.0 | U | 100 |
| 1,2,3,6,7,8-HxCDD | 0.1 | NotFnd | <0.90 | 0.90 | U | 100 |
| 1,2,3,7,8,9-HxCDD | 0.1 | NotFnd | <0.98 | 0.98 | U | 100 |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | NotFnd | <2.5 | 2.5 | U | 100 |
| OCDD | 0.0003 | 37.24 | 30.6 | 4.3 | M,J | 200 |
| 2,3,7,8-TCDF | 0.1 | NotFnd | <2.1 | 2.1 | U | 20 |
| 1,2,3,7,8-PeCDF | 0.03 | NotFnd | <1.0 | 1.0 | U | 100 |
| 2,3,4,7,8-PeCDF | 0.3 | NotFnd | <0.93 | 0.93 | U | 100 |
| 1,2,3,4,7,8-HxCDF | 0.1 | NotFnd | <0.85 | 0.85 | U | 100 |
| 1,2,3,6,7,8-HxCDF | 0.1 | NotFnd | <0.81 | 0.81 | U | 100 |
| 2,3,4,6,7,8-HxCDF | 0.1 | NotFnd | <0.89 | 0.89 | U | 100 |
| 1,2,3,7,8,9-HxCDF | 0.1 | NotFnd | <1.0 | 1.0 | U | 100 |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | NotFnd | <2.2 | 2.2 | U | 100 |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | NotFnd | <2.9 | 2.9 | U | 100 |
| OCDF | 0.0003 | NotFnd | <3.0 | 3.0 | U | 200 |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|----|-------|--------|
| 37C14-2,3,7,8-TCDD | 0 | NS | |
| 13C12-1,2,3,4,7,8-HxCDD | 0 | NS | |
| 13C12-2,3,4,7,8-PeCDF | 0 | NS | |
| 13C12-1,2,3,4,7,8-HxCDF | 0 | NS | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 0 | NS | |

| Extraction Standards | pg | Conc. | EDL |
|---------------------------|------|-------|-----------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.63 | 68 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.05 | 69 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.20 | 68 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.79 | 73 25-130 |
| 13C12-OCDD | 8000 | 37.23 | 73 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.72 | 70 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.06 | 72 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.71 | 72 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.23 | 77 25-130 |

| Cleanup Standard | pg | Conc. | EDL |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.47 | 70 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg | | |
|------------------------|---------|-------------|-----------|---|-----|
| Total-TCDD | 0 | <2.9 | 2.9 | U | 20 |
| Total-PeCDD | 0 | <1.7 | 1.7 | U | 100 |
| Total-HxCDD | 0 | <1.0 | 1.0 | U | 100 |
| Total-HpCDD | 0 | <2.5 | 2.5 | U | 100 |
| Total-TCDF | 0 | <2.1 | 2.1 | U | 20 |
| Total-PeCDF | 0 | <1.0 | 1.0 | U | 100 |
| Total-HxCDF | 0 | <1.0 | 1.0 | U | 100 |
| Total-HpCDF | 0 | <2.9 | 2.9 | U | 100 |

| | |
|--|-----------|
| Toxic Equivalency - (WHO 2005) | pg |
| Lower Bound PCDD/F TEQ (WHO 2005) | 0.00918 |
| Mid Point PCDD/F TEQ (WHO 2005) | 2.93 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 5.85 |

| | |
|------|--|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. |
| TEF | Indicates the Toxic Equivalency Factor |
| M | Indicates that a peak has been manually integrated. |
| U | Indicates that this compound was not detected above the EDL. |
| J | Indicates that a target analyte was detected below the calibrated range. |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure |
| NS | Indicates that this standards has not been added. |

ALS Life Sciences

Laboratory Method Blank Analysis Report

| | | | | | |
|--------------------|---------------------|------------------|-----------|--------|---|
| Sample Name | Method Blank | Sampling Date | n/a | | |
| ALS Sample ID | WG3463499-4 | Extraction Date | 23-Dec-21 | | Approved: <i>N Ashtari</i> --e-signature-- 28-Jan-2021 |
| Analysis Method | TO9A | Sample Size | 1 | Sample | |
| Analysis Type | Blank | Percent Moisture | n/a | | |
| Sample Matrix | REAGENT | Split Ratio | 4 | | |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210127A08 |
| Run Date | 27-Jan-21 21:58 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | pg |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | TEF (WHO 2005) | Ret. Time | Conc. pg | EDL pg Flags | EMPC pg | LQL |
|---------------------|-------------------|--------------|-------------|-----------------|------------|-----|
| 2,3,7,8-TCDD | 1 | NotFnd | <3.2 | 3.2 U | 20 | |
| 1,2,3,7,8-PeCDD | 1 | NotFnd | <1.7 | 1.7 U | 100 | |
| 1,2,3,4,7,8-HxCDD | 0.1 | NotFnd | <1.5 | 1.5 U | 100 | |
| 1,2,3,6,7,8-HxCDD | 0.1 | NotFnd | <1.3 | 1.3 U | 100 | |
| 1,2,3,7,8,9-HxCDD | 0.1 | NotFnd | <1.4 | 1.4 U | 100 | |
| 1,2,3,4,6,7,8-HpCDD | 0.01 | NotFnd | <2.2 | 2.2 U | 100 | |
| OCDD | 0.0003 | 37.25 | <10 | 4.2 M,J,R | 10 | 200 |
| 2,3,7,8-TCDF | 0.1 | NotFnd | <2.3 | 2.3 U | 20 | |
| 1,2,3,7,8-PeCDF | 0.03 | NotFnd | <1.0 | 1.0 U | 100 | |
| 2,3,4,7,8-PeCDF | 0.3 | NotFnd | <0.94 | 0.94 U | 100 | |
| 1,2,3,4,7,8-HxCDF | 0.1 | NotFnd | <0.93 | 0.93 U | 100 | |
| 1,2,3,6,7,8-HxCDF | 0.1 | NotFnd | <0.89 | 0.89 U | 100 | |
| 2,3,4,6,7,8-HxCDF | 0.1 | NotFnd | <0.98 | 0.98 U | 100 | |
| 1,2,3,7,8,9-HxCDF | 0.1 | NotFnd | <1.1 | 1.1 U | 100 | |
| 1,2,3,4,6,7,8-HpCDF | 0.01 | NotFnd | <0.85 | 0.85 U | 100 | |
| 1,2,3,4,7,8,9-HpCDF | 0.01 | NotFnd | <1.1 | 1.1 U | 100 | |
| OCDF | 0.0003 | NotFnd | <3.7 | 3.7 U | 200 | |

| Field Spike Standards | pg | % Rec | Limits |
|---------------------------|----|-------|--------|
| 37C14-2,3,7,8-TCDD | 0 | NS | |
| 13C12-1,2,3,4,7,8-HxCDD | 0 | NS | |
| 13C12-2,3,4,7,8-PeCDF | 0 | NS | |
| 13C12-1,2,3,4,7,8-HxCDF | 0 | NS | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 0 | NS | |

| Extraction Standards | pg | Conc. | EDL |
|---------------------------|------|-------|-----------|
| 13C12-2,3,7,8-TCDD | 4000 | 27.63 | 53 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.05 | 67 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.21 | 67 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.79 | 72 25-130 |
| 13C12-OCDD | 8000 | 37.24 | 69 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.72 | 60 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.07 | 63 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.72 | 71 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.24 | 78 25-130 |

| Cleanup Standard | pg | Conc. | EDL |
|-------------------------|------|-------|-----------|
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.47 | 61 40-130 |

| Homologue Group Totals | # peaks | Conc. pg | EDL pg | |
|------------------------|---------|-------------|-----------|-----|
| Total-TCDD | 0 | <3.2 | 3.2 U | 20 |
| Total-PeCDD | 0 | <1.7 | 1.7 U | 100 |
| Total-HxCDD | 0 | <1.5 | 1.5 U | 100 |
| Total-HpCDD | 0 | <2.2 | 2.2 U | 100 |
| Total-TCDF | 0 | <2.3 | 2.3 U | 20 |
| Total-PeCDF | 0 | <1.0 | 1.0 U | 100 |
| Total-HxCDF | 0 | <1.1 | 1.1 U | 100 |
| Total-HpCDF | 0 | <1.1 | 1.1 U | 100 |

| Toxic Equivalency - (WHO 2005) | pg |
|-----------------------------------|------|
| Lower Bound PCDD/F TEQ (WHO 2005) | 0.00 |
| Mid Point PCDD/F TEQ (WHO 2005) | 3.15 |
| Upper Bound PCDD/F TEQ (WHO 2005) | 6.30 |

| | |
|------|--|
| EDL | Indicates the Estimated Detection Limit, based on the measured background noise for this target in this sample. |
| TEF | Indicates the Toxic Equivalency Factor TEQ Indicates the Toxic Equivalency |
| M | Indicates that a peak has been manually integrated. |
| U | Indicates that this compound was not detected above the EDL. |
| | |
| J | Indicates that a target analyte was detected below the calibrated range. |
| R | Indicates that the ion abundance ratio for this compound did not meet the acceptance criterion. |
| | |
| LQL | Lower Quantification Limit, based on the lowest calibration level corrected for sample size, splits and dilutions. |
| EMPC | Estimated Maximum Possible Concentration - elevated detection limit due to interference or positive id criterion failure |
| NS | Indicates that this standards has not been added. |

ALS Life Sciences

Laboratory Control Sample Analysis Report

| | | | | | |
|--------------------|----------------------------------|------------------|-----------|--------|-------------------------------|
| Sample Name | Laboratory Control Sample | Sampling Date | n/a | | |
| ALS Sample ID | WG3463499-2 | Extraction Date | 23-Dec-21 | | |
| Analysis Method | T09A | Sample Size | 1 | Sample | Approved: <i>N Ashtari</i> |
| Analysis Type | LCS | Percent Moisture | n/a | | --e-signature-- |
| Sample Matrix | MEDIA | Split Ratio | 4 | | 28-Jan-2021 |

| | |
|------------------------|------------------------|
| Run Information | Run 1 |
| Filename | 7-210117A16 |
| Run Date | 17-Jan-21 18:09 |
| Final Volume | 10 uL |
| Dilution Factor | 1 |
| Analysis Units | % |
| Instrument - Column | HRMS-7 DB5MSUS0710421H |

| Target Analytes | pg | Ret. Time | % Rec | Limits Flags |
|------------------------------|-----------|--------------|--------------|-----------------|
| 2,3,7,8-TCDD | 400 | 27.71 | 90 | 70-130 |
| 1,2,3,7,8-PeCDD | 2000 | 32.09 | 105 | 70-130 |
| 1,2,3,4,7,8-HxCDD | 2000 | 34.17 | 102 | 70-130 |
| 1,2,3,6,7,8-HxCDD | 2000 | 34.23 | 100 | 70-130 |
| 1,2,3,7,8,9-HxCDD | 2000 | 34.35 | 103 | 70-130 |
| 1,2,3,4,6,7,8-HpCDD | 2000 | 35.82 | 98 | 70-130 |
| OCDD | 4000 | 37.27 | 110 | 70-130 |
| 2,3,7,8-TCDF | 400 | 26.80 | 91 | 70-130 |
| 1,2,3,7,8-PeCDF | 2000 | 31.12 | 90 | 70-130 |
| 2,3,4,7,8-PeCDF | 2000 | 31.87 | 82 | 70-130 |
| 1,2,3,4,7,8-HxCDF | 2000 | 33.68 | 90 | 70-130 |
| 1,2,3,6,7,8-HxCDF | 2000 | 33.75 | 95 | 70-130 |
| 2,3,4,6,7,8-HxCDF | 2000 | 34.08 | 86 | 70-130 |
| 1,2,3,7,8,9-HxCDF | 2000 | 34.50 | 86 | 70-130 |
| 1,2,3,4,6,7,8-HpCDF | 2000 | 35.26 | 89 | 70-130 |
| 1,2,3,4,7,8,9-HpCDF | 2000 | 36.06 | 92 | 70-130 |
| OCDF | 4000 | 37.36 | 79 | 70-130 |
| Field Spike Standards | pg | | % Rec | Limits |
| 37Cl4-2,3,7,8-TCDD | 0 | | NS | |
| 13C12-1,2,3,4,7,8-HxCDD | 0 | | NS | |
| 13C12-2,3,4,7,8-PeCDF | 0 | | NS | |
| 13C12-1,2,3,4,7,8-HxCDF | 0 | | NS | |
| 13C12-1,2,3,4,7,8,9-HpCDF | 0 | | NS | |
| Extraction Standards | | | | |
| 13C12-2,3,7,8-TCDD | 4000 | 27.68 | 52 | 40-130 |
| 13C12-1,2,3,7,8-PeCDD | 4000 | 32.08 | 53 | 40-130 |
| 13C12-1,2,3,6,7,8-HxCDD | 4000 | 34.22 | 56 | 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDD | 4000 | 35.81 | 54 | 25-130 |
| 13C12-OCDD | 8000 | 37.27 | 59 | 25-130 |
| 13C12-2,3,7,8-TCDF | 4000 | 26.77 | 56 | 40-130 |
| 13C12-1,2,3,7,8-PeCDF | 4000 | 31.10 | 50 | 40-130 |
| 13C12-1,2,3,6,7,8-HxCDF | 4000 | 33.73 | 56 | 40-130 |
| 13C12-1,2,3,4,6,7,8-HpCDF | 4000 | 35.25 | 53 | 25-130 |
| Cleanup Standard | pg | | | |
| 13C12-1,2,3,7,8,9-HxCDF | 4000 | 34.49 | 54 | 40-130 |

NS Indicates that this standards has not been added.

SVOC DATA PACKAGE

SECTION 6: INTERNAL RECORDS

Including:

- Prep Logs
- Independent calculation checks
- Others as listed below:

Batch ID:

WG3463499

Batch ID: WG3463499

DX Native Standard:

| Sample I.D. | Volume (ul) | (Checkmark) Spiked |
|-------------|-------------|-----------------------|
| WG3463499-2 | 40 | ✓ |
| WG3463499-3 | 40 | ✓ |

PCB Native Standard:

| Sample I.D. | Volume (ul) | (Checkmark) Spiked |
|-------------|-------------|-----------------------|
| WG3463499-2 | 40 | ✓ |
| WG3463499-3 | 40 | ✓ |

DX Cleanup Standard:

| Sample I.D. | Volume (ul) | (Checkmark) Spiked |
|-------------|-------------|-----------------------|
| WG3463499-1 | 20 | ✓ |
| WG3463499-2 | 20 | ✓ |
| WG3463499-3 | N/A | N/A |
| WG3463499-4 | 20 | ✓ |
| L2541483-1 | 20 | ✓ |
| L2541483-2 | 20 | ✓ |
| L2541483-3 | 20 | ✓ |
| L2541483-4 | 20 | ✓ |
| L2541483-5 | 20 | ✓ |
| | 20 | |
| | 20 | |
| | | |
| | | |
| | | |
| | | |
| | | |
| | | |
| | | |

PCB Cleanup Standard:

| Sample I.D. | Volume (ul) | (Checkmark) Spiked |
|-------------|-------------|-----------------------|
| WG3463499-1 | 20 | ✓ |
| WG3463499-2 | 20 | ✓ |
| WG3463499-3 | N/A | N/A |
| WG3463499-4 | 20 | ✓ |
| L2541483-1 | 20 | ✓ |
| L2541483-2 | 20 | ✓ |
| L2541483-3 | 20 | ✓ |
| L2541483-4 | 20 | ✓ |
| L2541483-5 | 20 | ✓ |
| | 20 | |
| | 20 | |
| | | |
| | | |
| | | |
| | | |
| | | |
| | | |

Syringe ID: 322

Standard: 1613B-NS#3-030B

Date & Initials: 23-Dec-20 SS

Syringe ID: 394

Standard: 1668A-NS#1-040D

Date & Initials: 23-Dec-20 SS

Syringe ID: 357

Standard: M23-CL#1-037A

Date & Initials: 28-Dec-20 SS/Bm

Correct Syringe Obtained: SS/Bm
Chemist's Initials

Correct Standard Obtained: SS/Bm
Chemist's Initials

Correct Technique Followed: SS/Bm
Chemist's Initials

Syringe ID: 395

Standard: 1668A-CL#2-038J

Date & Initials: 28-Dec-20 SS/Bm

Correct Syringe Obtained: SS/Bm
Chemist's Initials

Correct Standard Obtained: SS/Bm
Chemist's Initials

Correct Technique Followed: SS/Bm
Chemist's Initials

Batch ID: WG3463499

Reagent Lot Numbers:

| Reagent | Lot# | Manufacturer |
|-----------------------|---------------------------|--------------|
| Acetone | 105642 | |
| Hexane | 105523 | |
| DCM | 105939 | |
| Toluene | 105795 | |
| Nonane | ORG-WAKONON- 056 | |
| 1:1 DCM:HEX | ORG-DH2- 662 | |
| Sodium Sulphate | ORG-SSU- 2462, 2461 | |
| Acid Silica | ORG-ASI- 9524, 9525, 9526 | |
| Neutral Silica | ORG-NSI- 2463 | |
| Alumina | ORG-ALU- 491 | |
| 1% Deactivated Silica | ORG-2%DAS- - | |
| Chromacarb | ORG-CC- 255 | |

Cornoil

073

Batch ID: WG3463499

Procedure:

This batchsheet is a guideline only. Please see test procedure for complete set of instructions.

Extraction:

- For MB and LCS you **must** use blank media - if not available see your Team Lead

XAD+Filter+Front Half Rinse

- Place a layer of pre-cleaned glasswool with 1cm of sodium sulphate or a preclened thimble into soxhlet

- Place the PUFFS into the soxhlet body or precleaned thimble

- Spike with Extraction Standard (plus Native for LCS and ENI).

- Soxhlet extract in DCM for 16 hours (check with team lead or supervisor)

Rotovap:

- Rotovap and reduce to ~2mL.

- Transfer to a calibrated c-tube (marked at 1ml, 2ml) with 3x2ml hexane

- Mix well then quantitatively spilt the extract **1/2 DX/PCB 1/2 Archive**

Batch ID: WG3463499

DX/PCB:

- Perform Acid Silica column \
- Solvent Exchange (reduce to **~50ul**, bulk back up to 1ml Hexane, vortex well. ✓
- Perform Alumina Column:
 - Pre-elute the Alumina Column with 7ml Hexane ✓
 - Place F1 c-tube under the column, then load the sample with 3x1ml Hexane rinses ✓
 - F1 (Archive) 1mL Hexane ✓
 - F2 (DX/PCB) 14mL 1:1 DCM:Hexane

-Split Alumina F2 1/2 PCB 1/2 DX

Micro-Vial:

PCB:

- Blow down to ~1/2ml
- Vortex **very** well.
- Transfer every last drop to a micro-vial (Marked at 20uL with nonane).
- Blow down to the line
- Spike PCB Injection Standard, cap and vortex. **FV=25ul**

DX:

- Solvent Exchange to Hexane (Reduce to Just Dry then bulk back up to 1ml Hexane)

- ChromaCarb: - 4cm of well-packed chroma-carb.

- Pre-elute Carbon with 5ml Hexane

- Transfer with 3x1ml Hexane

- F1 = **10ml** 1:1 DCM:Hexane (Archive)

- After dripping has stopped Invert Column.

- F2 = 14ml Toluene (DX and PCB)

- After the column has stopped dripping reduce the **F2** portion down to ~1/2ml.

- Vortex well, then transfer to a micro-vial without rinses.

- Blow the micro-vial down to just-dry.

- Spike with Injection Standard, Cap the micro-vial, and Vortex. **FV=10ul**

Batch ID: WG3463499

Comments:

NOTE: Label and Save All Columns including Acid Silica Columns

Approval of Deviation from Standard Method

Procedure does deviate from Standard Method. **Approved (Supervisor/Manager):** _____

(Batch Writer): _____

Procedure does deviate from Standard Method. **Approved (Supervisor/Manager):** _____

| | | | | | |
|-----------------------|-----------|------------------|---------------|-----------|--|
| WG3463499 | | Prep Analyst: | | | |
| PUFS - M23/1668A (HR) | | Date: | | | |
| | Very Good | Meets Method Req | Some Outliers | Very Poor | Comments / Was spl/batch sent for rework? Why? |
| MB | | | | | |
| LCS | | | | | |
| DUP | | | | | |
| ES rec | | | | | |

ALS Life Sciences

Sample Calculation Report

CS3 RRF Check

Approved:

N Ashtari
 --e-signature--
 28-Jan-2021

$$\text{RRF} = \frac{\text{Response of 2,3,7,8-TCDD}}{\text{Response of 13C12-2,3,7,8-TCDD}} \times \frac{\text{Concentration of 13C12-2,3,7,8-TCDD}}{\text{Concentration of TCDD}}$$

$$\text{RRF} = \frac{167775.70}{1460254.80} \times \frac{100}{10}$$

Calculated Value
Value from TargetLynx

$$= 1.149 \qquad 1.149$$

Calculation of OCDD amount in L2541483-1

$$\text{pg} = \frac{\text{Response of OCDD}}{\text{Response of 13C12-OCDD}} \times \frac{\text{pg of 13C12-OCDD spiked}}{\text{Mean RRF} * \text{Sample Size}}$$

$$\text{pg} = \frac{47999.5}{343802.4} \times \frac{8000}{0.96 * 1.00} = 1159 \qquad 1159$$

Calculation of 13C12-2,3,7,8-TCDD Recovery in L2541483-1

$$\% \text{ Recovery} = \frac{\text{Response of 13C12-2,3,7,8-TCDD}}{\text{Response of 13C12-1,2,3,4-TCDD}} \times \frac{\text{pg of 13C12-1,2,3,4-TCDD spiked} * 100}{\text{Mean RRF} * \text{Amount Spiked}}$$

$$\% \text{ Recovery} = \frac{1211291.2}{1547309.8} \times \frac{4000 * 100}{1.11 * 4000} = 71 \qquad 71 \%$$

SVOC DATA PACKAGE

SECTION 7: SHIPPING/RECEIVING DOCUMENTS

Including:

- Airbills
- Chain-of-Custody Records
- Sample Log-in Sheet(s) - where applicable
- Others as listed below:



Chain of Custody (COC) / Analytical Request Form



Composite WO: L2541483

COC Number: 17-792310

L2541477-COFC

Page of

Canada Toll Free: 1 800 668 9878

www.alsglobal.com

| | | | | | | | | | | | | | | | | | | | | | | | |
|--|---|---|----------------------------------|---|--------------------|---|-------|----------------------|--|--|--|---|--|--|--|------------------|--|--|--|---------|--|--|--|
| Report To Contact and company name below will appear on the final report. | | Report Format / Distribution | | Select Service Level Below - Contact your AM to confirm all E&P TATs (surcharges may apply) | | | | | | | | | | | | | | | | | | | |
| Company: <u>Farallon Consulting</u> | | Select Report Format: <input checked="" type="checkbox"/> PDF <input checked="" type="checkbox"/> EXCEL <input checked="" type="checkbox"/> EDD (DIGITAL) | | Regular [R] <input checked="" type="checkbox"/> Standard TAT if received by 3 pm - business days - no surcharges apply | | | | | | | | | | | | | | | | | | | |
| Contact: <u>Amber Bailey</u> | | Quality Control (QC) Report with Report <input type="checkbox"/> YES <input type="checkbox"/> NO | | Priority (Business Days) | | EMERGENCY | | | | | | | | | | | | | | | | | |
| Phone: <u>206-735-6178</u> | | <input type="checkbox"/> Compare Results to Criteria on Report - provide details below if box checked | | 4 day [P4-20%] <input type="checkbox"/> | | 1 Business day [E - 100%] <input type="checkbox"/> | | | | | | | | | | | | | | | | | |
| Company address below will appear on the final report | | Select Distribution: <input type="checkbox"/> EMAIL <input type="checkbox"/> MAIL <input type="checkbox"/> FAX | | 3 day [P3-25%] <input type="checkbox"/> | | Same Day, Weekend or Statutory holiday [E2 - 200%] (Laboratory opening fees may apply) <input type="checkbox"/> | | | | | | | | | | | | | | | | | |
| Street: <u>975 5th AVE NW</u> | | Email 1 or Fax: <u>ambert@farallonconsulting.com</u> | | Date and Time Required for all E&P TATs: dd-mmm-yy hh:mm | | | | | | | | | | | | | | | | | | | |
| City/Province: <u>ISSAQUAH WA</u> | | Email 2: <u>spatterson@farallonconsulting.com</u> | | For tests that can not be performed according to the service level selected, you will be contacted. | | | | | | | | | | | | | | | | | | | |
| Postal Code: <u>98059</u> | | Email 3: | | Analysis Request | | | | | | | | | | | | | | | | | | | |
| Invoice To | | Invoice Distribution | | Indicate Filtered (F), Preserved (P) or Filtered and Preserved (F/P) below | | | | | | | | | | | | | | | | | | | |
| Same as Report To <input checked="" type="checkbox"/> YES <input type="checkbox"/> NO | | Select Invoice Distribution: <input checked="" type="checkbox"/> EMAIL <input type="checkbox"/> MAIL <input type="checkbox"/> FAX | | <table border="1"> <tr><td colspan="4">NUMBER OF CONTAINERS</td></tr> <tr><td colspan="4">9855 EPA Method</td></tr> <tr><td colspan="4">EPA Method 8240A</td></tr> <tr><td colspan="4">Dioxins</td></tr> </table> | | | | NUMBER OF CONTAINERS | | | | 9855 EPA Method | | | | EPA Method 8240A | | | | Dioxins | | | |
| NUMBER OF CONTAINERS | | | | | | | | | | | | | | | | | | | | | | | |
| 9855 EPA Method | | | | | | | | | | | | | | | | | | | | | | | |
| EPA Method 8240A | | | | | | | | | | | | | | | | | | | | | | | |
| Dioxins | | | | | | | | | | | | | | | | | | | | | | | |
| Copy of Invoice with Report <input checked="" type="checkbox"/> YES <input type="checkbox"/> NO | | Email 1 or Fax: <u>Amber@farallonconsulting.com</u> | | | | | | | | | | | | | | | | | | | | | |
| Company: | | Email 2: | | | | | | | | | | | | | | | | | | | | | |
| Contact: | | Email 3: | | | | | | | | | | | | | | | | | | | | | |
| Project Information | | Oil and Gas Required Fields (client use) | | <table border="1"> <tr><td colspan="4">SAMPLES ON HOLD</td></tr> <tr><td colspan="4">SUSPECTED HAZARD (see Special Instructions)</td></tr> </table> | | | | SAMPLES ON HOLD | | | | SUSPECTED HAZARD (see Special Instructions) | | | | | | | | | | | |
| SAMPLES ON HOLD | | | | | | | | | | | | | | | | | | | | | | | |
| SUSPECTED HAZARD (see Special Instructions) | | | | | | | | | | | | | | | | | | | | | | | |
| ALS Account # / Quote #: | | AFE/Cost Center: | | | | | | | | | | | | | | | | | | | | | |
| Job #: | | Major/Minor Code: | | | | | | | | | | | | | | | | | | | | | |
| PO / AFE: | | Requisitioner: | | | | | | | | | | | | | | | | | | | | | |
| LSD: | | Location: | | | | | | | | | | | | | | | | | | | | | |
| ALS Lab Work Order # (lab use only): | | ALS Contact: | | Sampler: | | | | | | | | | | | | | | | | | | | |
| ALS Sample # (lab use only) | Sample Identification and/or Coordinates (This description will appear on the report) | Date (dd-mmm-yy) | Time (hh:mm) | Sample Type | | | | | | | | | | | | | | | | | | | |
| 1 | L2497422-8-1 | 12/15/20 | 1241 | AW | 1 | X | X | | | | | | | | | | | | | | | | |
| 2 | L2497422-1-2 | | 1250 | | 1 | X | X | | | | | | | | | | | | | | | | |
| 3 | L2497422-9-3 | | 1324 | | 1 | X | X | | | | | | | | | | | | | | | | |
| 4 | L2497422-7-4 | | 1334 | | 1 | X | X | | | | | | | | | | | | | | | | |
| 5 | L2497422-10-5 | | 1343 | | 1 | X | X | | | | | | | | | | | | | | | | |
| Drinking Water (DW) Samples¹ (client use) | | Special Instructions / Specify Criteria to add on report by clicking on the drop-down list below (electronic COC only) | | SAMPLE CONDITION AS RECEIVED (lab use only) | | | | | | | | | | | | | | | | | | | |
| Are samples taken from a Regulated DW System? <input type="checkbox"/> YES <input type="checkbox"/> NO | | Analyze samples - composite samples submitted on 10/15, 11/17, and 12/15. | | Frozen <input type="checkbox"/> SIF Observations Yes <input type="checkbox"/> No <input checked="" type="checkbox"/> | | | | | | | | | | | | | | | | | | | |
| Are samples for human consumption/use? <input type="checkbox"/> YES <input type="checkbox"/> NO | | | | Ice Packs <input checked="" type="checkbox"/> Ice Cubes <input type="checkbox"/> Custody seal intact Yes <input checked="" type="checkbox"/> No <input type="checkbox"/> | | | | | | | | | | | | | | | | | | | |
| | | | | Cooling Initiated <input checked="" type="checkbox"/> | | | | | | | | | | | | | | | | | | | |
| | | | | INITIAL COOLER TEMPERATURES °C: 5.4°C | | | | | | | | | | | | | | | | | | | |
| | | | | FINAL COOLER TEMPERATURES °C: | | | | | | | | | | | | | | | | | | | |
| SHIPMENT RELEASE (client use) | | INITIAL SHIPMENT RECEPTION (lab use only) | | FINAL SHIPMENT RECEPTION (lab use only) | | | | | | | | | | | | | | | | | | | |
| Released by: <u>[Signature]</u> | Date: <u>12/15/20</u> | Time: <u>1500</u> | Received by: <u>PARAN BURTON</u> | Date: <u>13-DEC-2020</u> | Time: <u>13:10</u> | Received by: | Date: | | | | | | | | | | | | | | | | |

REFER TO BACK PAGE FOR ALS LOCATIONS AND SAMPLING INFORMATION

WHITE - LABORATORY COPY YELLOW - CLIENT COPY

ALS Canada Ltd.

L2541483 DX DPKG 210129

60 of 65

1. If any water samples are taken from a Regulated Drinking Water (DW) System, please submit using an Authorized DW COC form.

Sample Receiving Log

| Date/Time Received | Client ID | Number/Description of Containers | Temp. on Receipt* | Condition of Samples, Courier & Tracking Information | Receiver's Initials | Date/Time Login Completed | Submission ID | Sample ID Range |
|----------------------|-----------|----------------------------------|-------------------|--|---------------------|---------------------------|---------------|-----------------|
| | | | | | | | | |
| 21-Oct-2020 11:00 | FARALLON | 5 x PUFs | 10.8°C | Good FedEx 7714 6810 5736 | Mg | 21-Oct-2020 14:45 | L2579524 | -1-5 |

*Temperatures were recorded using : VWR Traceable dedicated I.R. gun (model 36934-178 SN 192108143)
 Other (specify): _____

Sample Receiving Log

| Date/Time Received | Client ID | Number/Description of Containers | Temp. on Receipt* | Condition of Samples, Courier & Tracking Information | Receiver's Initials | Date/Time Login Completed | Submission ID | Sample ID Range |
|----------------------|-----------|----------------------------------|-------------------|--|---------------------|---------------------------|---------------|-----------------|
| | | | | | | | | |
| 18-NOV-2026 11:45 | FARALLON | 5 x PUFFS | 5.9°C | Good FedEx 770 6531 8900 | Mg | 18-NOV-2026 14:08 | L2530845 | -1-5 |
| | | | | | | | | |

*Temperatures were recorded using : VWR Traceable dedicated I.R. gun (model 36934-178 SN 192108143)
 Other (specify): _____

Sample Receiving Log

| Date/Time Received | Client ID | Number/Description of Containers | Temp. on Receipt* | Condition of Samples, Courier & Tracking Information | Receiver's Initials | Date/Time Login Completed | Submission ID | Sample ID Range |
|----------------------|-----------|----------------------------------|-------------------|--|---------------------|---------------------------|----------------------|-----------------|
| | | | | | | | | |
| 16-Dec-2020 13:10 | FARALLON | 5 x PUFs | 5.4°C | Good FedEx 7718 6371 7805 | MB | 17-Dec-2020 12:45 | L2541477 L2541483 | -1-5 -1-5 |

*Temperatures were recorded using : VWR Traceable dedicated I.R. gun (model 36934-178 SN 192108143)
 Other (specify): _____

From: [Claire Kocharakkal](#)
To: [Amber Bailey](#); [Breanne Dusureault](#)
Subject: RE: [EXTERNAL] - COC from 12/15/20
Date: Friday, January 8, 2021 12:40:54 PM
Attachments: [image003.png](#)
[image004.png](#)
[image005.png](#)

Hello Amber,

We can certainly look into this! Breanne will be your account manager going forward and she will be happy to check in on this and respond.

Thank you!

Claire Kocharakkal
Account Manager, Environmental
Canada



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E +1 905 331 4567
claire.kocharakkal@alsglobal.com
1435 Norjohn Court Unit 1
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From: Amber Bailey [mailto:abailey@farallonconsulting.com]
Sent: Friday, January 08, 2021 12:37 PM
To: Claire Kocharakkal <claire.kocharakkal@ALSGlobal.com>
Subject: [EXTERNAL] - COC from 12/15/20
Importance: High

CAUTION: This email originated from outside of ALS. Do not click links or open attachments unless you recognize the sender and are sure content is relevant to you.

Claire,

Can you double check on something for me? I think I wrote the incorrect sample label IDs on the COC from the 12/15 sample submission.

If so please verify that the following samples were actually the ones submitted.

Site 1: L2527465-3
Site 2: L2527465- 2
Site 3: L2527465- 5
Site 4: L2527465- 4
Site 5: L2527465- 1

If these are correct please correct on the COC to:

L2527465-3-1
L2527465- 2- 2
L2527465- 5- 3
L2527465- 4- 4
L2527465- 1- 5

I apologize for the mix up.

Thank you,

Amber Bailey, Project Environmental Scientist

Farallon Consulting, L.L.C. | 975 5th Avenue Northwest | Issaquah, Washington 98027
abailey@farallonconsulting.com | Direct: (425) 295-0811 Cell: (206) 735-6178



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