



Report of Analysis

Client: Nobis Group

Project: Raymark Superfund Site

Client Sample ID: PIBLK-PS028848.D

Lab Sample ID: I.BLK-PS028848.D

Analytical Method: SW8151A

Sample Wt/Vol: 1000 Units: mL

Soil Aliquot Vol: uL

Extraction Type:

PS028848.D

GPC Factor: 1.0 PH:

Prep Method: SW3510C

File ID/Qc Batch: Dilution:

Prep Date

Date Analyzed

Date Collected:

Date Received:

SDG No.:

Matrix:

% Solid:

Final Vol:

Injection Volume:

Test:

12/30/24

12/30/24

P5306

10000

Herbicide Group1

WATER

Prep Batch ID

Decanted:

иL

12/30/24 ps123024

| CAS Number | Parameter | Conc. | Qualifier | MDL | LOD LOQ/CRQL | | Units |
|------------|-------------------|--------|-----------|----------|--------------|--------|----------|
| TARGETS | | | | | | | |
| 1918-00-9 | DICAMBA | 0.0015 | U | 0.00042 | 0.0015 | 0.0020 | mg/L |
| 75-99-0 | DALAPON | 0.0015 | U | 0.0011 | 0.0015 | 0.0020 | mg/L |
| 120-36-5 | DICHLORPROP | 0.0015 | U | 0.00043 | 0.0015 | 0.0020 | mg/L |
| 94-75-7 | 2,4-D | 0.0015 | U | 0.00049 | 0.0015 | 0.0020 | mg/L |
| 93-72-1 | 2,4,5-TP (Silvex) | 0.0015 | U | 0.00045 | 0.0015 | 0.0020 | mg/L |
| 93-76-5 | 2,4,5-T | 0.0015 | U | 0.00050 | 0.0015 | 0.0020 | mg/L |
| 94-82-6 | 2,4-DB | 0.0015 | U | 0.00057 | 0.0015 | 0.0020 | mg/L |
| 88-85-7 | DINOSEB | 0.0015 | U | 0.00055 | 0.0015 | 0.0020 | mg/L |
| SURROGATES | | | | | | | |
| 19719-28-9 | 2,4-DCAA | 599 | | 32 - 138 | | 120% | SPK: 500 |

Comments:

U = Not Detected

LOO = Limit of Quantitation

MDL = Method Detection Limit

LOD = Limit of Detection

E = Value Exceeds Calibration Range

P = Indicates >25% difference for detected concentrations between the two GC columns

Q = indicates LCS control criteria did not meet requirements

M = MS/MSD acceptance criteria did not meet requirements

J = Estimated Value

B = Analyte Found in Associated Method Blank

N = Presumptive Evidence of a Compound

* = Values outside of QC limits

D = Dilution

S = Indicates estimated value where valid five-point calibration was not performed prior to analyte detection in sample.

() = Laboratory InHouse Limit