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UCD CSN Technical Information #801B

CSN Data Processing

Chemical Speciation Network Air Quality Research Center University of California, Davis

> July 31, 2019 Version 1.2

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DOCUMENT HISTORY

Date Modified	Initials	Section/s Modified	Brief Description of Modifications
11/30/18	NJS	All	Reworded for clarity, equations added, and updated name changes.
7/31/2019	DEY, KAG	1,3,7,8,9,10	Changed process for carbon and ion analysis pathways, and wording changes for clarity.

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1. PURPOSE AND APPLICABILITY

The subject of this technical information (TI) document is processing the sampling and analytical data from the Chemical Speciation Network (CSN). The raw operational information from field sampling is combined with laboratory analytical results to generate concentrations, uncertainties, and method detection limits.

2. SUMMARY OF THE METHOD

The University of California, Davis (UCD) analyst will use functions in the *crocker* software package to calculate final results and post them to the UCD CSN database. The analyst will also review any output messages for errors.

3. **DEFINITIONS**

- **crocker:** A custom software package in the R language that contains the data processing code used to produce, check, and post the final results.
- **CSN database:** A SQL Server database that is the central warehouse of CSN preliminary and final data at UCD.
- MDL: Method detection limit.
- **RCM:** Reconstructed Mass.

4. HEALTH AND SAFETY WARNINGS

Not applicable.

5. CAUTIONS

Not applicable.

6. INTERFERENCES

Not applicable.

7. PERSONNEL QUALIFICATIONS, DUTIES, AND TRAINING

The UCD Air Quality Research Center (AQRC) Data & Reporting Group staff assigned to tasks described in this document have advanced training in database programming and database management.

8. PROCEDURAL STEPS

Data processing is performed using the *crocker* R package, which is developed and maintained by UCD specifically for data processing, monitoring, and validation of the

CSN data. Data processing is performed by the UCD Data & Reporting Group on monthly batches of data (a calendar month of sample start dates). Processing occurs in two steps.

- 1. Calculate concentrations, uncertainties, and MDLs, and check for duplicates (Section 8.1).
- 2. Post results to the UCD CSN database (Section 8.2).

8.1 Calculate concentrations, uncertainties, and MDLs, and check for duplicates

Laboratory results for ions and carbon fractions are stored in the database as mass per filter and elements are stored as mass per cm². The *csn_calculate* methods in the *crocker* package combine per filter analysis results with filter operational data (e.g., flow rates) and corresponding blanks to calculate concentrations, uncertainty estimates, and MDLs. The details and specific equations are provided in Section 9.

To calculate values for all measured and derived parameters, the analyst should open an R environment (such as RStudio) and run the following command¹:

[month_data] <- crocker::csn_calculate_all([YYYY], [MM], 'production')

This command will calculate concentrations, uncertainties, and MDLs for all measured and derived parameters for the year ([YYYY]) and month ([MM]) and return them (in memory) to the variable [month_data]. The last argument in the command specifies that the calculations will use the "production" database (i.e., the CSN operational database).

Within the *csn_calculate_all* function, the concentrations, uncertainties, and MDLs are calculated for each of the three main groups of species (elements, ions, and carbon) as well as for calculated values such as sulfate, nitrate, soil, and OMC, and reconstructed mass. Following the calculations, a check for duplicates is performed and processing will cease if duplicates are identified. When duplicate check indicates a problem (which is typical), the analyst will investigate and resolve the issue. Duplicate issues can result from the same filter having more than one set of analysis results or because there appears to be more than one filter of the same filter type running on the same day at the same site. Once the issues are resolved, the *csn_calculate_all* command should be re-run.

8.2 Post results to CSN database

After all duplicates have been resolved, the results can be uploaded to the UCD CSN production database in preparation for Level 1 data validation. The operator will execute the following commands:

[post] <- crocker::csn_post_results([month_data], 'production', AnalysisQcCode = 1, comment = ['Initial Posting'], replacingId = NULL, replacingQcCode = NULL)

This command appends the processed data to the analysis.Results table in the UCD CSN production database as an analysis set. It also inserts a record into the analysis.ResultsSets table that provides summary information for this set, including the

¹ Text in [brackets] indicates values that can be changed by the user. Other values should be typed as written.

comment and the *AnalysisQcCode*. *AnalysisQcCode* = 1 is used for routine data. During level 1 validation, the data may be modified and $csn_calculate_all$ is run again and a new complete data set is posted to the database. The analyst should use the $csn_post_results$ command again, change the comment to record what the dataset relates to as well as changing the analysis QC code of the previously posted dataset(s) by including the data set ID of the previous posting (*replacingId*) and the analysis QC code (*replacingQcCode*) that should be associated with that data set.

9. DATA PROCESSING EQUATIONS

Section 9.1 through Section 9.4 present the equations used to calculate species concentrations and the associated uncertainties and MDLs. These calculations are performed by the *crocker* R package.

9.1 Ions

Ions are measured on the nylon filter by ion chromatography (IC). The ions measurement – chloride (Cl⁻), nitrate (NO₃⁻), sulfate (SO₄²⁻), ammonium (NH₄⁺), potassium (K⁺), and sodium (Na⁺) – are delivered as micrograms per filter in the ions data files from RTI. The ion concentration, MDL, and uncertainty in micrograms per cubic meter are calculated using Equations 1, 2, and 3 respectively.

$$C = \frac{A - B}{V}$$
 1

Where,

C = ambient concentration (μ g/m³)

A = mass measured on sample (μ g/filter)

B = artifact mass (μ g/filter) = monthly median of field blanks (FB)

 $V = sample volume (m^3)$

The MDL is calculated as the 95th percentile minus the median of monthly field blanks, with a minimum analytical value and dependency on sample volume (Equation 2). The MDL is reported with each sample concentration and varies per month of reported data and also with sample volume.

$$MDL = \frac{max \begin{cases} P_{95} - med\\ analytical MDL \end{cases}}{V}$$

Where,

 $P_{95} = 95$ th percentile of at least 50 field blanks for the matching month, including nearby months if necessary (µg/filter)

med = Median (50th percentile) of at least 50 field blanks for the matching month, including nearby months if necessary (μ g/filter)

analytical MDL = analytical MDL value as reported by the ion chromatography laboratory (μ g); this value is updated periodically per direction from the analytical laboratory

 $V = sample volume (m^3)$

Uncertainty is reported with each concentration and varies for each concentration reported (Equation 3).

$$\sigma(c) = \sqrt{(S_{FB})^2 + (fC)^2} \qquad 3$$

Where,

 S_{FB} = analytical uncertainty, calculated as (Equation 4).

$$S_{FB} = \frac{MDL}{1.6449}$$

 $C = ambient concentration (\mu g/m^3)$

f = fractional uncertainty (Equation 6), calculated using scaled relative differences (SRD; Equation 5) between sample pairs collected at CSN collocated sites using the most recent two years of data (currently, June 1, 2016 through May 31, 2018). If the total number of valid collocated pairs over the two-year period is less than 60, a value of 0.25 is adopted as the fractional uncertainty. The fractional uncertainty is updated annually (corresponding with delivery of January data).

scaled relative difference (srd) =
$$\frac{(collocated - routine)/\sqrt{2}}{(collocated - routine)/2}$$
 5

Fractional Uncertainty
$$(f) = \frac{(84th \ percentile \ of \ srd) - (16th \ percentile \ of \ srd)}{2}$$
 6

The analytical MDL and fractional uncertainty for each ion species is stored in the database and shown in Table 1.

Species	Analytical MDL	f
Chloride	0.104	0.122
Nitrate	0.067	0.108
Sulfate	0.123	0.085
Ammonium	0.019	0.209
Sodium	0.044	0.163
Potassium	0.587	0.179

Table 1. Analytical MDL and fractional uncertainty (f) for ions.

9.2 Carbon

Carbon is measured on the quartz filer by thermal optical analysis (TOA). Carbon measurements are reported from the laboratory as micrograms per filter. For the eight carbon species, the primary source of fractional uncertainty is the separation into temperature ranges. This may be associated with temperature regulation, but it may also be from the inherent variability of the species involved. The concentration, MDL, and uncertainty in micrograms per cubic meter for the carbon fractions and sums – OC1, OC2, OC3, OC4, OPTR, OPTT, EC1, EC2, EC3, OCTR, ECTR, and TCTC – are calculated both with and without an artifact correction using Equations 7 through 11. For the versions without artifact correction (e.g., OC1_raw, ECTR_raw), the B variable in Equation 7 is 0. For the values with artifact correction (e.g., OC1, ECTR), the B variable in Equation 7 is the monthly median mass loading of field blanks across the network.

$$C = \frac{A - B}{V}$$
 7

Where,

 $C = ambient concentration (\mu g/m^3)$

A = mass measured on sample (μ g/filter)

 $B = artifact mass (\mu g/filter) = monthly median of field blanks$

V =sample volume (m³)

The MDL is calculated as the 95th percentile minus the median of monthly field blanks, with a minimum analytical value and dependency on sample volume (Equation 8). The MDL is reported with each sample concentration and varies per month of reported data and also with sample volume.

$$MDL = \frac{max \begin{cases} P_{95} - med\\ analytical MDL \end{cases}}{V}$$

Where,

 $P_{95} = 95$ th percentile of at least 50 field blanks for the matching month, including nearby months if necessary (µg/filter)

med = median (50th percentile) of at least 50 field blanks for the matching month, including nearby months if necessary (μ g/filter)

analytical MDL = analytical MDL value as reported by the thermal optical analysis laboratory (μ g); this value is updated periodically per direction from the analytical laboratory.

Uncertainty is reported with each concentration and varies for each concentration reported (Equation 9).

$$\sigma(c) = \sqrt{(S_{FB})^2 + (fC)^2}$$

Where,

 S_{FB} = analytical uncertainty, calculated as (Equation 10).

$$S_{FB} = \frac{MDL}{1.6449}$$
10

C = ambient concentration (μ g/m³)

f = fractional uncertainty (Equation 12), calculated using scaled relative differences (SRD; Equation 11) between sample pairs collected at CSN collocated sites using the most recent two years of data (currently, June 1, 2016 through May 31, 2018). If the total number of valid collocated pairs over the two-year period is less than 60, a value of 0.25 is adopted as the fractional uncertainty. The fractional uncertainty is updated annually (corresponding with delivery of January data).

scaled relative difference (srd) =
$$\frac{(collocated - routine)/\sqrt{2}}{(collocated - routine)/2}$$
 11

Fractional Uncertainty
$$(f) = \frac{(84th \ percentile \ of \ srd) - (16th \ percentile \ of \ srd)}{2}$$
 12

The analytical MDL and fractional uncertainty for each carbon species is stored in the database and shown in Table 2.

Species	Analytical MDL	f
OC1	0.17	0.0.335
OC2	1.05	0.125
OC3	0.42	0.151
OC4	0.26	0.194
OPTR	0.23	0.377
OPTT	0.23	0.254
OCTR	1.43	0.103
OCTT	1.48	0.094
EC1	0.22	0.152
EC2	0.22	0.320
EC3	0.06	0.25
ECTR	0.37	0.169
ECTT	0.37	0.183
ТСТС	1.48	0.25

Table 2. Analytical MDL and fractional uncertainty (f) for artifact corrected carbon species.

9.3 Elements

Elements are measured on the PTFE filter using X-ray fluorescence (XRF). Because the XRF instrument reports areal densities, concentrations (*C*) are calculated using Equation 13.

$$C = \frac{AD \cdot a - L}{V}$$
 13

Where,

AD = areal density ($\mu g/cm^2$)

L = median mass loading of at least 50 field blanks for the matching month, including nearby months if necessary (µg/filter)

a = sample deposit area (cm²)

 $V = sample volume (m^3)$

The MDL is calculated as the 95th percentile minus the median of monthly field blanks, with a minimum analytical value and dependency on sample volume (Equation 14). The MDL is reported with each sample concentration and varies per month of reported data and also with sample volume.

$$MDL = \frac{max \begin{cases} P_{95} - med\\ analytical MDL \end{cases}}{V}$$
14

Where,

 $P_{95} = 95$ th percentile of at least 50 field blanks for the matching month, including nearby months if necessary (µg/filter)

med = median (50th percentile) of at least 50 field blanks for the matching month, including nearby months if necessary (μ g/filter)

analytical MDL = analytical MDL value as reported by the x-ray fluorescence laboratory (μ g); this value is updated periodically per direction from the analytical laboratory.

Uncertainty is reported with each concentration and varies for each concentration reported (Equation 15).

$$\sigma(c) = \sqrt{(S_{FB})^2 + (fC)^2}$$
¹⁵

Where,

 S_{FB} = analytical uncertainty, calculated as (Equation 16).

$$S_{FB} = \frac{MDL}{1.6449} \tag{16}$$

 $C = ambient concentration (\mu g/m^3)$

f = fractional uncertainty (Equation 18), calculated using scaled relative differences (SRD; Equation 17) between sample pairs collected at CSN collocated sites using the most recent two years of data (currently, June 1, 2016 through May 31, 2018). If the total number of valid collocated pairs over the two-year period is less than 60, a value of 0.25 is adopted as the fractional uncertainty. The fractional uncertainty is updated annually (corresponding with delivery of January data).

scaled relative difference (srd) =
$$\frac{(collocated - routine)/\sqrt{2}}{(collocated - routine)/2}$$
 17

Fractional Uncertainty
$$(f) = \frac{(84th \ percentile \ of \ srd) - (16th \ percentile \ of \ srd)}{2}$$
 18

The analytical MDL and fractional uncertainty for each carbon species is stored in the database and shown in Table 2.

Parameter	Analytical MDL	f
Ag	0.159	0.25
Al	0.312	0.132
As	0.018	0.25
Ba	0.776	0.25
Br	0.044	0.25
Ca	0.054	0.115
Cd	0.153	0.25
Ce	0.924	0.25
Cl	0.02	0.364
Со	0.032	0.25
Cr	0.026	0.25
Cs	0.521	0.25
Cu	0.11	0.25
Fe	0.17	0.144
In	0.369	0.25
K	0.021	0.081
Mg	0.397	0.25
Mn	0.062	0.25
Na	0.858	0.185
Ni	0.018	0.25
Р	0.02	0.25
Pb	0.118	0.25
Rb	0.086	0.25
S	0.036	0.056
Sb	0.376	0.25
Se	0.051	0.25
Si	0.133	0.141
Sn	0.473	0.25
Sr	0.07	0.25
Ti	0.031	0.18
V	0.013	0.25
Zn	0.03	0.11
Zr	0.348	0.25

Table 3. Analytical MDL and fractional uncertainty (f) for elemental species.

9.4 Reconstructed Mass

Reconstructed mass (RCM; Equation 19) is a composite variable estimated from chemical speciation measurements, and can be compared to gravimetric measurements to asses overall data quality. Gravimetric analysis is not routinely performed using CSN filters; thus, for comparison purposes data from collocated continuous monitors (available from AirNow Tech) are used for validation.

$$RCM = NHSO + NHNO + Soil + 1.8 Chloride + ECTR + OMC$$
 19

Where,

NHSO = ammonium sulfate concentration (see 9.4.1) NHNO = ammonium nitrate concentration (see 9.4.2) Soil = soil concentration (see 9.4.3) Chloride = chloride concentration as measured by IC (Section 9.1)

ECTR = elemental carbon concentration by TOA (Section 9.2)

OMC = concentration of organic mass by carbon (see 9.4.4)

For all of the terms in Equation 19, zero is substituted for negative values. The MDL for RCM is zero, and uncertainty is calculated as combination of the individual uncertainties (Equation 20).

$$\sigma_{RCM} = \sqrt{\sigma_{NHSO}^2 + \sigma_{NHNO}^2 + \sigma_{Soil}^2 + (1.8 \sigma_{Chloride})^2 + \sigma_{ECTR}^2 + \sigma_{OMC}^2}$$
 20

For Equation 20, each uncertainty term must be at least zero.

9.4.1 Ammonium Sulfate (NHSO)

Sulfur is predominantly present as sulfate in the atmosphere, generally as ammonium sulfate $(NH_4)_2SO_4$, though also as ammonium bisulfate $(NH_4)HSO_4$, sulfuric acid H₂SO₄, gypsum CaSO₄·2H₂O, and, in marine areas, sodium sulfate NaSO₄. In many cases, the particle will include associated water, but this is omitted from the calculation. All sulfur is assumed to be present as ammonium sulfate (NHSO), and the concentration, MDL, and uncertainty as:

$$C_{NHSO} = C_S * 4.125 \tag{21}$$

$$MDL_{NHSO} = MDL_s * 4.125$$

$$\sigma_{NSHO} = \sigma_S * 4.125 \tag{23}$$

9.4.2 Ammonium Nitrate (NHNO)

This is the total dry concentration associated with nitrate, assuming 100% neutralization by ammonium. The concentration, MDL, and uncertainty are derived from the nitrate ion measurement and are calculated as:

$$C_{NHNO} = C_{nitrate} * 1.29$$

$$MDL_{NHn0} = MDL_{nitrate} * 1.29$$
 25

$$\sigma_{NSNO} = \sigma_{nitrate} * 1.29$$

9.4.3 Soil

The soil component consists of the sum of the predominantly soil elements measured by XRF, multiplied by a coefficient to account for oxygen for the normal oxide forms (Al₂O₃, SiO₂, CaO, K₂O, FeO, Fe₂O₃, TiO₂), and augmented by a factor to account for other compounds not included in the calculation, such as MgO, Na₂O, water, and CO₂. The following assumptions are made:

• Fe is split equally between FeO (oxide factor of 1.29) and Fe_2O_3 (oxide factor of 1.43), giving an overall Fe oxide factor of 1.36.

• Fine K has a non-soil component from smoke. Based on the K/Fe ratio for average sediment (*Handbook of Chemistry and Physics*) of 0.6, 0.6*Fe is used as a surrogate for soil K and then add the oxide factor for K₂O (oxide factor of 1.2) to get a total Fe factor of 0.72*Fe (0.6*1.2) for the potassium oxide in soil. This increases the factor for Fe from 1.36 to 2.08.

• The oxide forms of the soil elements account for 86% of average sediment; in order to obtain the total mass associated with soil, the final factors are divided by 0.86 (*Handbook of Chemistry and Physics*). The final equations for fine soil concentration, MDL, and uncertainty are:

$$C_{soil} = 2.2 Al + 2.49 Si + 1.63 Ca + 2.42 Fe + 1.94 Ti$$
27

$$MDL_{soil} = max(2.2 MDL_{Al}, 2.49 MDL_{Si}, 1.63 MDL_{Ca}, 2.42 MDL_{Fe}, 1.94 MDL_{Ti})$$
28

$$\sigma_{soil} = \sqrt{(2.2 \,\sigma_{Al})^2 + (2.49 \,\sigma_{Si})^2 + (1.63 \,\sigma_{Ca})^2 + (2.42 \,\sigma_{Fe})^2 + (1.94 \,\sigma_{Ti})^2}$$
²⁹

For Equations 30-32, zero substitution is applied to any negative terms.

9.4.4 Organic Mass by Carbon (OMC)

To determine the total amount of organic mass associated with organic carbon, the ratio of organic mass to organic carbon is assumed to be 1.4.

$$C_{OMC} = 1.4 * C_{OCTR}$$
 30

$$MDL_{OMC} = 1.4 * MDL_{OCTR}$$
 31

$$\sigma_{OMC} = 1.4 * \sigma_{OCTR}$$
 32

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10. DATA PROCESSING CODE

This section describes the flow of data through the data processing code used to calculate concentration, MDL, and uncertainty for all CSN parameters. Figure 1 outlines the flow of data from the filter and analysis specific database tables to final results. The wrapper function *csn_calculate_all* is the only function executed directly by the analyst (see Section 8.1); *csn_calculate_all* in turn calls several functions sequentially to calculate first measured and then derived concentrations. Source code for the functions shown in Figure 1 is stored in the UCD source repository.

Figure 1. Flow diagram of the processing code in *crocker::csn_calculate_all*. The raw sampling and analysis data are inputs to the process. Rectangles represent R functions and arrows indicate inputs and outputs.



11. EQUIPMENT AND SUPPLIES

The associated hardware and software used for CSN data ingest are described in the associated UCD CSN SOP #801: Processing & Validating Raw Data.

12. QUALITY ASSURANCE AND QUALITY CONTROL

Software bugs and data management issues are tracked through JIRA tracking software. All users have access to our internal JIRA website and can submit, track, and comment on bug reports.

13. REFERENCES

Not Applicable.