

# Synoptic CB: Porewater DIC

August 2023 Samples

2025-10-25

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```
##Setup - Change things here & write any notes
```

```
#identify section  
cat("Setup Information")
```

```
## Setup Information
```

```
##### Run information - PLEASE CHANGE  
Date_Run = "08/18/23" #Date that instrument was run  
Run_by = "Stephanie J. Wilson" #Instrument user  
Script_run_by = "Stephanie J. Wilson" #Code user  
run_notes = " " #any notes from the run  
samples <- c("GCW", "GWI", "MSM", "SWH") #whatever identifies your samples within the same names  
samples_pattern <- paste(samples, collapse = "|")  
  #samples_pattern <- "GCW" #use this instead of the line above if you have only one site code  
chks_name = "Chk_Std_" #what did you name your check standards?  
crm_name = "CRM|crm" #what did you name your CRMS?  
  
##### File Names - PLEASE CHANGE  
#file path and name for raw summary data file  
raw_file_name = c("Raw Data/TOCTN_COMPASS_Synoptic_DIC_202308_1.txt",  
                  "Raw Data/TOCTN_COMPASS_Synoptic_DIC_202308_2.txt")  
  
#file path and name for raw all peaks file  
# raw_allpeaks_name = c("Raw Data/COMPASS_SynopticCB_PW_DIC_202308_1_allpeaks.txt",  
#                       "Raw Data/COMPASS_SynopticCB_PW_DIC_202308_2_allpeaks.txt")  
  
#file path and name of processed data file  
processed_file_name = "Processed Data/COMPASS_SynopticCB_PW_Processed_DIC_202308.csv"  
  
##### Log Files - PLEASE CHECK  
#downloaded metadata csv - downloaded from Google drive as csv for this year  
Raw_Metadata = "Raw Data/COMPASS_SynopticCB_PW_SampleLog_2023.csv"  
  
#qaqc log file path for this year  
#Log_path = "Raw Data/COMPASS_Synoptic_DIC_QAQClog_2023.csv"
```

```
##Set Up Code
```

```
##Read in metadata and create similar sample IDs for matching to samples
```

## 0.1 Import Data Functions

## 0.2 Import Sample Data

```
## Import Sample Data
```

```
## New names:  
## New names:  
## * ' ' -> '...14'
```

```
## # A tibble: 6 x 3
##   sample_name      ic_raw run_datetime
##   <chr>          <dbl> <chr>
## 1 GCW_202308_TR_LysB_10cm    28.1 8/19/2023 9:50:53 AM
## 2 GCW_202308_WC_LysA_10cm   162. 8/19/2023 10:08:13 AM
## 3 GCW_202308_WC_LysA_20cm   148. 8/19/2023 10:26:42 AM
## 4 GCW_202308_WC_LysA_45cm   170. 8/19/2023 10:44:08 AM
## 5 GCW_202308_WC_LysB_10cm   198. 8/19/2023 11:00:44 AM
## 6 GCW_202308_WC_LysB_20cm   165. 8/19/2023 11:18:50 AM
```

### 0.3 Assessing Standard Curves - assessed on the instrument manually

### 0.4 CRM Check - Don't run chunk if no CRMs run

```
## Assess the CRMs
```

```
## New names:
## New names:
## * ' ' -> '...14'
```

```
## [1] "IC crm has a % Difference <25% of expected - PROCEED"
```

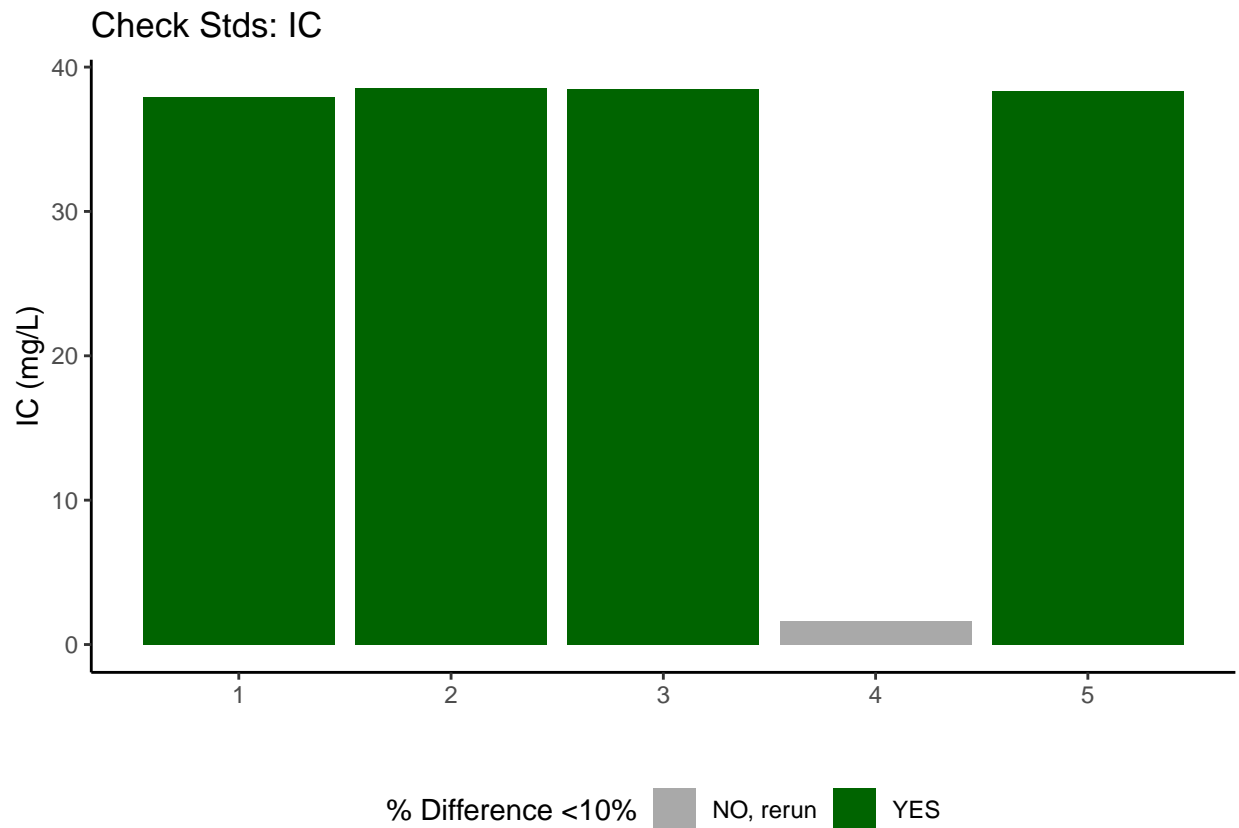
```
## Run mean = 23.1575
```

```
## Expected = 22.19
```

### 0.5 Assess Check Standards

```
## Assess the Check Standards
```

```
## New names:
## New names:
## * ' ' -> '...14'
```



```
## [1] ">60% of IC Check Standards are within range of expected concentration"
```

## 0.6 Assess Blanks

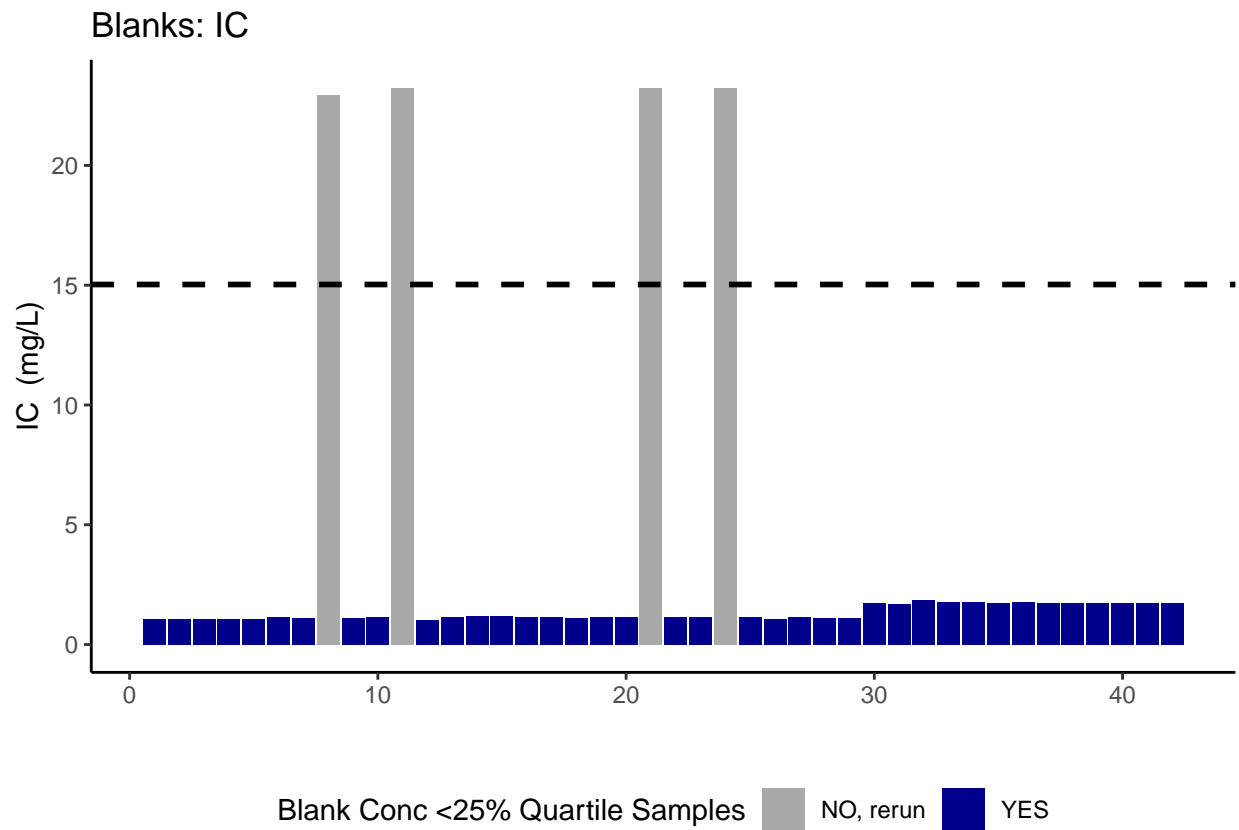
```
## Assess Blanks
```

```
## New names:
```

```
## New names:
```

```
## * ' ' -> '...14'
```

```
## [1] ">60% of Carbon Blank concentrations are lower 25% quartile of samples"
```

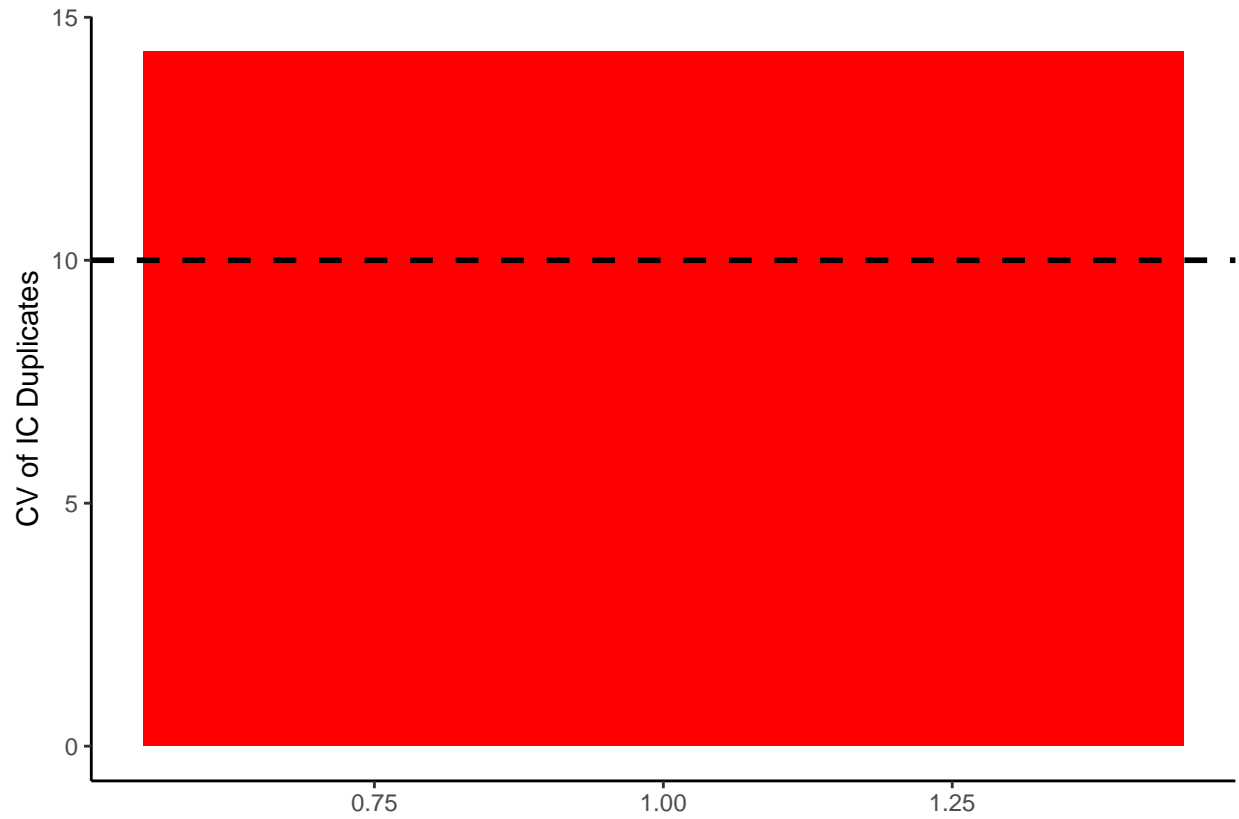


```
## carbon blanks:
```

```
## [1] 3.413595
```

## 0.7 Assess Duplicates

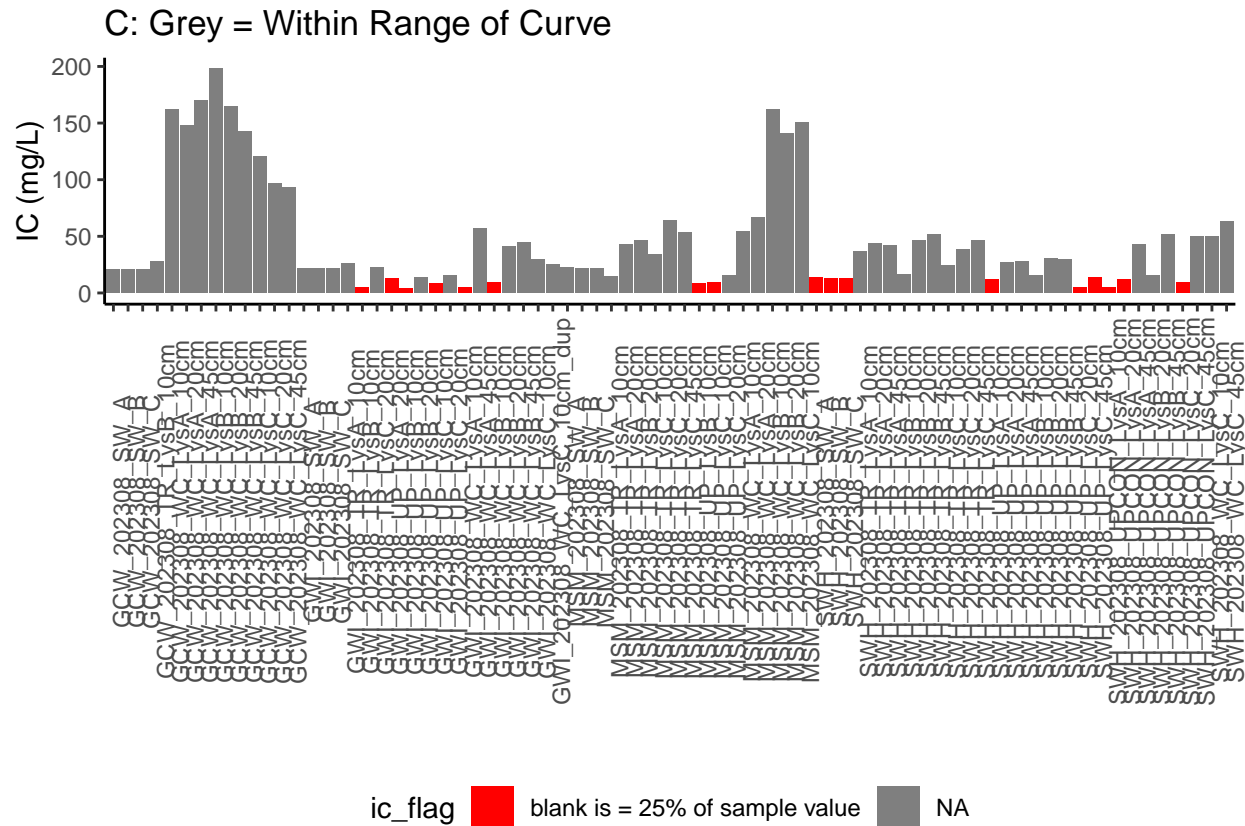
```
## Assess Duplicates
```



```
## [1] "<60% of Carbon Duplicates have a CV <10% - REASSESS"
```

## 0.8 Sample Flagging - Are samples Within the range of the curve?

```
## Sample Flagging
```



## 0.9 Visualize Data by Plot

```
## Visualize Data
```

```
## Warning in rbind(c("GCW", "202308", "TR", "LysB", "10cm"), c("GCW", "202308", :
## number of columns of result is not a multiple of vector length (arg 1)
```

