

# COMPASS\_Synoptic\_TGW\_2023: November Samples that got Skipped

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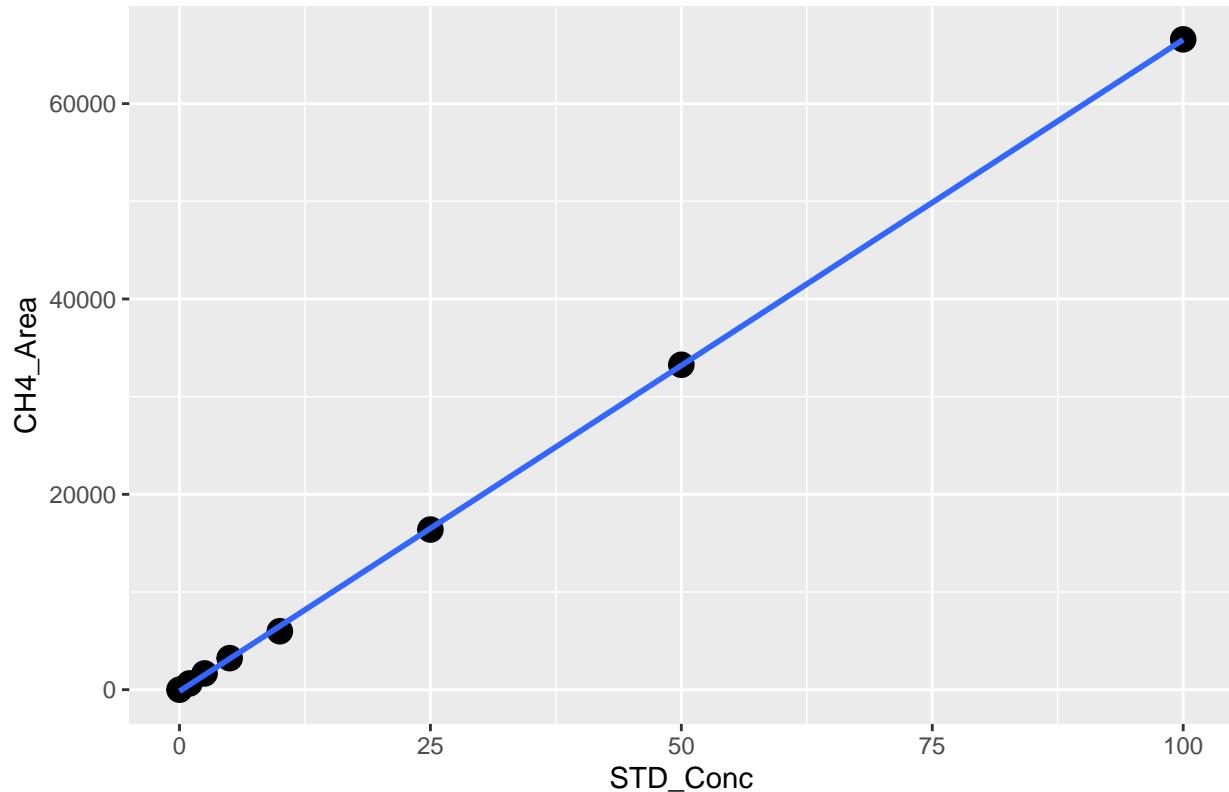
##Set Up

**Read in first data file and assess standard curves**

```
##      Machine User Run_Date Sample_Type Type1 Sample_Year Sample_Month
## 1 Varian GC Wegner 20240311     Blank    Blank        2024       <NA>
## 2 Varian GC Wegner 20240311 STD_CH4 STD_CH4        2024       <NA>
## 3 Varian GC Wegner 20240311 STD_CO2 STD_CO2        2024       <NA>
## 4 Varian GC Wegner 20240311 STD_CH4 STD_CH4        2024       <NA>
## 5 Varian GC Wegner 20240311 STD_CH4 STD_CH4        2024       <NA>
## 6 Varian GC Wegner 20240311 STD_CH4 STD_CH4        2024       <NA>
##               Sample_ID Dilution_Factor STD_Conc CO2_Area CH4_Area Field.Notes
## 1             Blank_0                 1     0.0     8170        0       NA
## 2 Blank_0_repeatforCH4                 1     0.0     8170        0       NA
## 3 Blank_0_repeatforCO2                 1     0.0     8170        0       NA
## 4           STD_1ppm_CH4                 1     1.0    25120       642       NA
## 5           STD_2.5ppm_CH4                1     2.5    69200      1664       NA
## 6           STD_5ppm_CH4                 1     5.0   141017      3228       NA
##   Lab.Notes
## 1
## 2
## 3
## 4
## 5
## 6

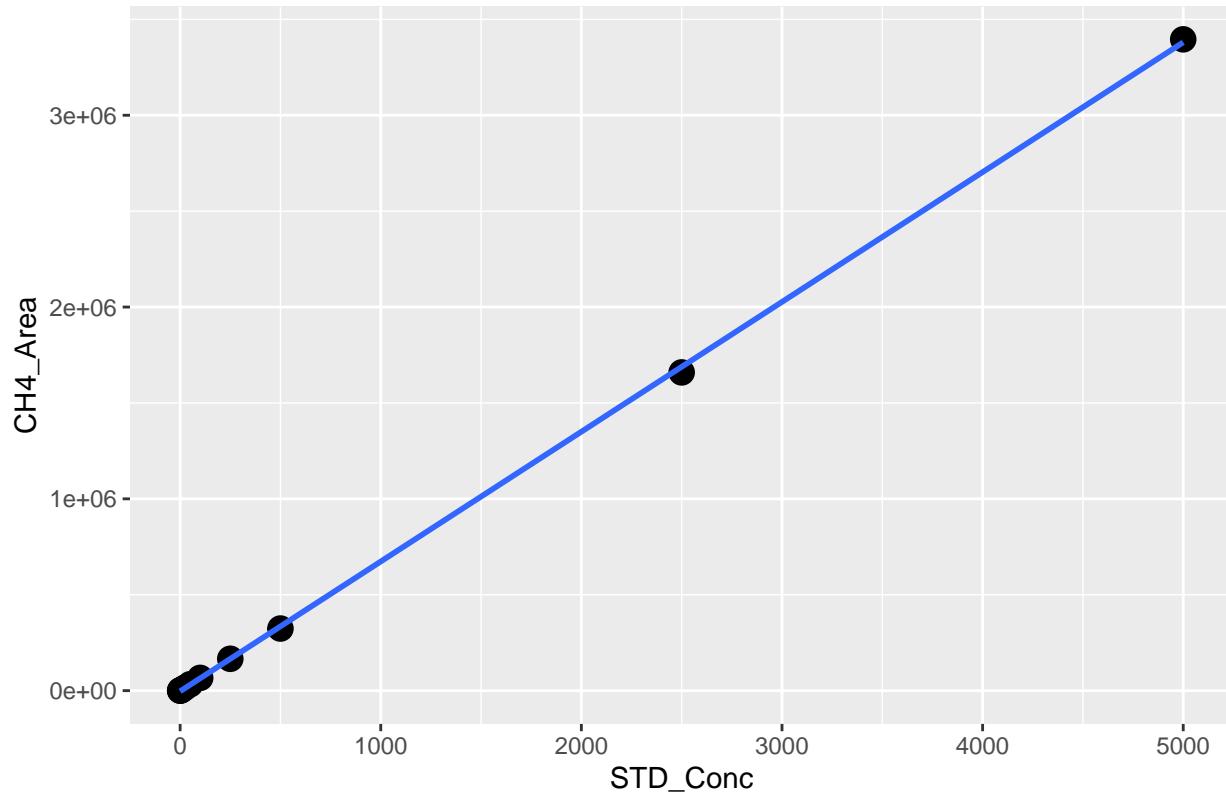
## `geom_smooth()` using formula = 'y ~ x'
```

## CH4 LOW Std Curve



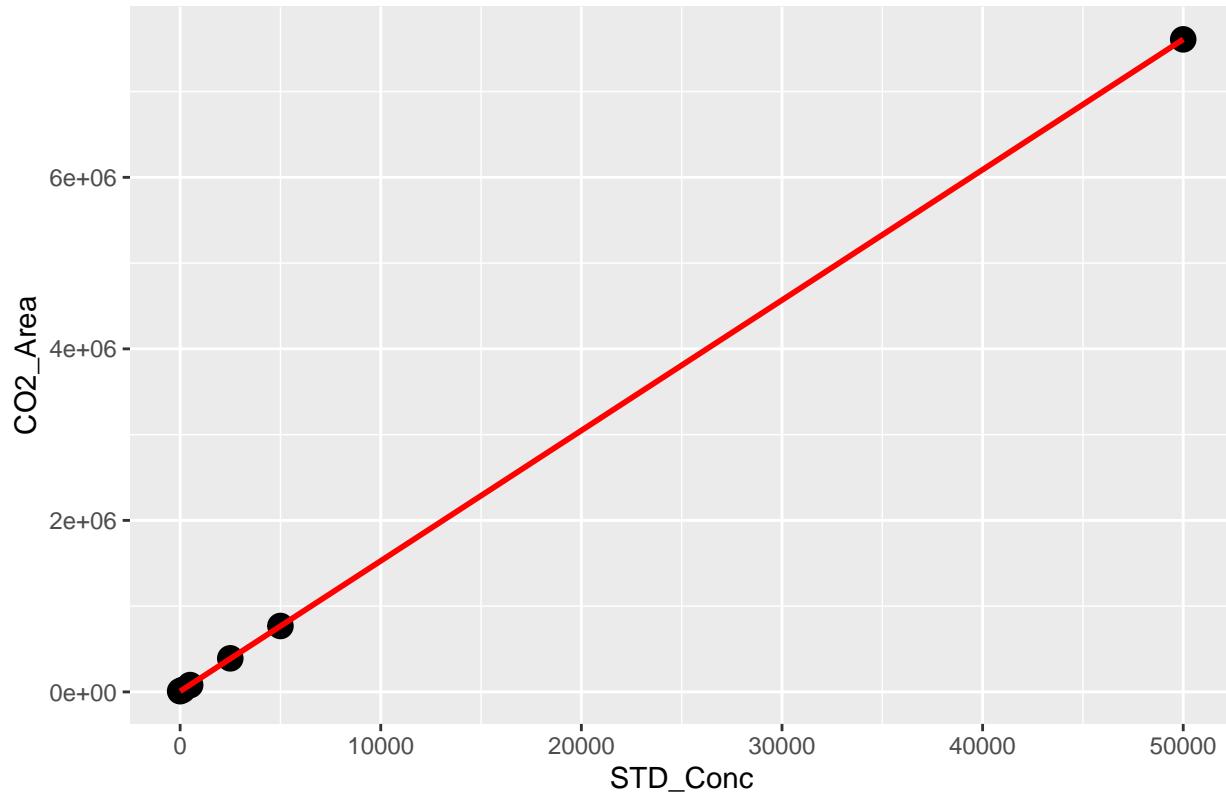
```
##  
## Call:  
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -520.10    -2.44    66.29   146.31   166.10  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (Intercept)           -166.101    108.922  -1.525  0.178  
## stds_ch4_low$STD_Conc  667.120     2.676 249.328 2.81e-13 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 247.8 on 6 degrees of freedom  
## Multiple R-squared:  0.9999, Adjusted R-squared:  0.9999  
## F-statistic: 6.216e+04 on 1 and 6 DF,  p-value: 2.809e-13  
  
## 'geom_smooth()' using formula = 'y ~ x'
```

## CH4 HIGH Std Curve



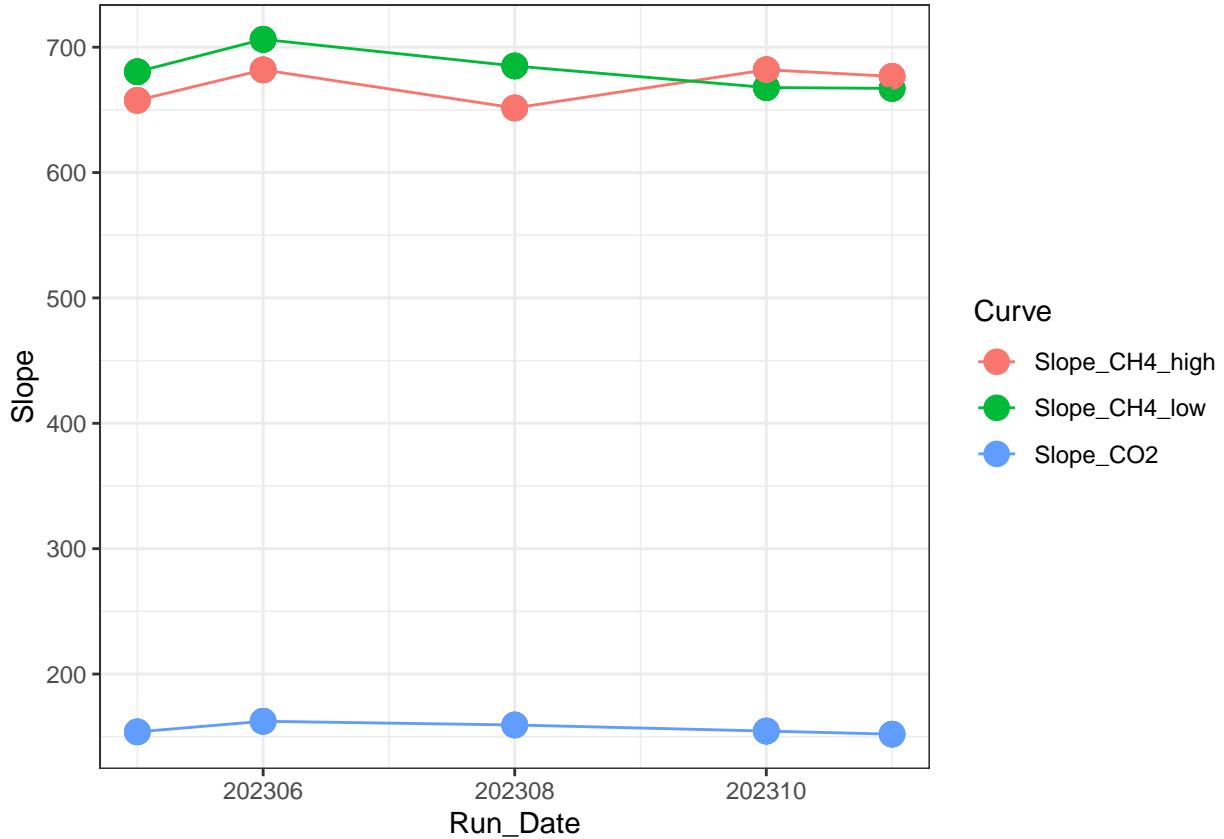
```
##  
## Call:  
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -29185    1847    2900    3416   15587  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (Intercept)           -3448.891   3644.622  -0.946   0.366  
## stds_ch4$STD_Conc    676.747     2.247 301.204 <2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 11380 on 10 degrees of freedom  
## Multiple R-squared:  0.9999, Adjusted R-squared:  0.9999  
## F-statistic: 9.072e+04 on 1 and 10 DF,  p-value: < 2.2e-16  
  
## 'geom_smooth()' using formula = 'y ~ x'
```

## CO2 Std Curve



```
##
## Call:
## lm(formula = stds_co2$CO2_Area ~ stds_co2$STD_Conc)
##
## Residuals:
##      1       2       3       4       5       6 
## 1296.2 -3146.3 -3365.7  4181.7  1335.2 -301.1 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 6.874e+03 1.505e+03  4.567  0.0103 *  
## stds_co2$STD_Conc 1.520e+02 7.327e-02 2075.176 3.24e-13 *** 
## ---      
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1 
##
## Residual standard error: 3251 on 4 degrees of freedom
## Multiple R-squared:      1, Adjusted R-squared:      1 
## F-statistic: 4.306e+06 on 1 and 4 DF,  p-value: 3.235e-13 

##      X          Curve      R2      Slope Intercept Run_Date
## 1 1 Slope_CH4_low 0.9984062 680.4131 -473.3666 202305
## 2 2 Slope_CH4_high 0.9998709 657.6131 3119.1378 202305
## 3 3 Slope_CO2 0.9999897 153.8752 9651.4986 202305
## 4 4 Slope_CH4_low 0.9984062 680.4131 -473.3666 202305
## 5 5 Slope_CH4_high 0.9998709 657.6131 3119.1378 202305
## 6 6 Slope_CO2 0.9999897 153.8752 9651.4986 202305
```



Now calculate the CH4 & CO2 concentrations in ppm

```
#head(raw)

#pull out methane standards
Samples <- raw %>%
  filter(!str_detect(Sample_Type, "STD_CH4")) %>%
  filter(!str_detect(Sample_Type, "STD_CO2")) %>%
  filter(!str_detect(Sample_Type, "Blank")) %>%
  filter(!str_detect(Sample_Type, "Chk_STD")) %>%
  filter(!str_detect(Sample_Type, "CHKSTD")) %>%
  filter(!str_detect(Sample_Type, "CHK_STD")) %>%
  filter(!str_detect(Sample_ID, "Blank")) %>%
  filter(!str_detect(Sample_ID, "CHKSTD")) %>%
  filter(!str_detect(Sample_Type, "NA"))

#head(Samples)

#Now flag any areas that are above the 100ppm area for CH4
Samples$CH4_Curve <- ifelse(Samples$CH4_Area >71000, "High", "Low")
#head(Samples)

#Calculate CH4 concentrations in ppm
Samples$CH4_Conc_ppm <- ifelse(Samples$CH4_Area >71000, (Samples$CH4_Area - Slope_CH4_high$Intercept)/Slope_CH4_high$Slope, (Samples$CH4_Area - Slope_CH4_low$Intercept)/Slope_CH4_low$Slope)
```

```

#Calculate CO2 concentrations in ppm
Samples$CO2_Conc_ppm <- ((Samples$CO2_Area-Slope_CO2$Intercept)/Slope_CO2$Slope)

#head(Samples)

#####make flags for any dilutions needed
#highest CH4 standard = 10000
#highest CO2 standard = 50000

Samples$CH4_Flag <- ifelse(Samples$CH4_Conc_ppm >10000, "Needs Dilution", "Within Range")
Samples$CO2_Flag <- ifelse(Samples$CO2_Conc_ppm >50000, "Needs Dilution", "Within Range")
#head(Samples)

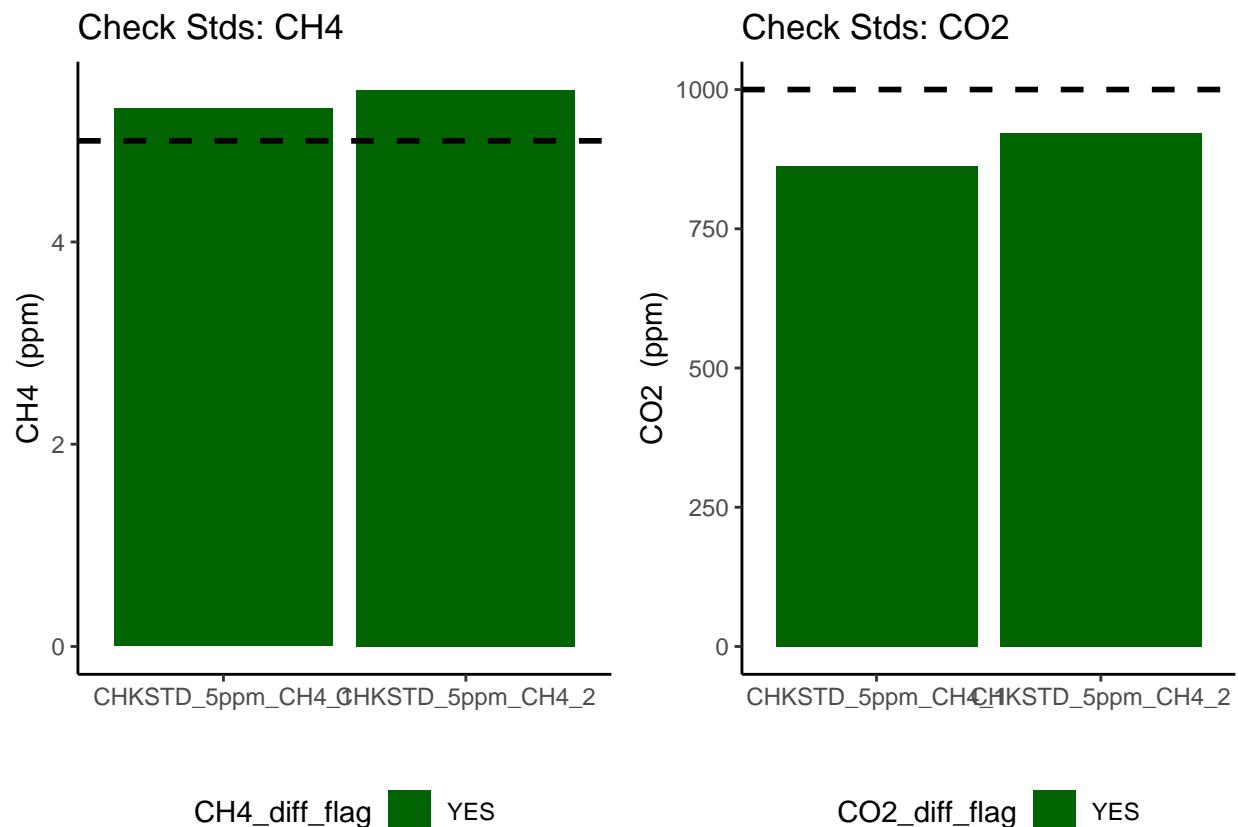
```

## Check the Check Standards

```

## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.
## i Please use 'linewidth' instead.
## This warning is displayed once every 8 hours.
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was
## generated.

```



## Dilution correct samples

```
#multiply the concentration by the dilution factor
Samples$CH4_Conc_ppm_dilcorr <- (Samples$CH4_Conc_ppm * Samples$Dilution_Factor)

Samples$CO2_Conc_ppm_dilcorr <- (Samples$CO2_Conc_ppm * Samples$Dilution_Factor)

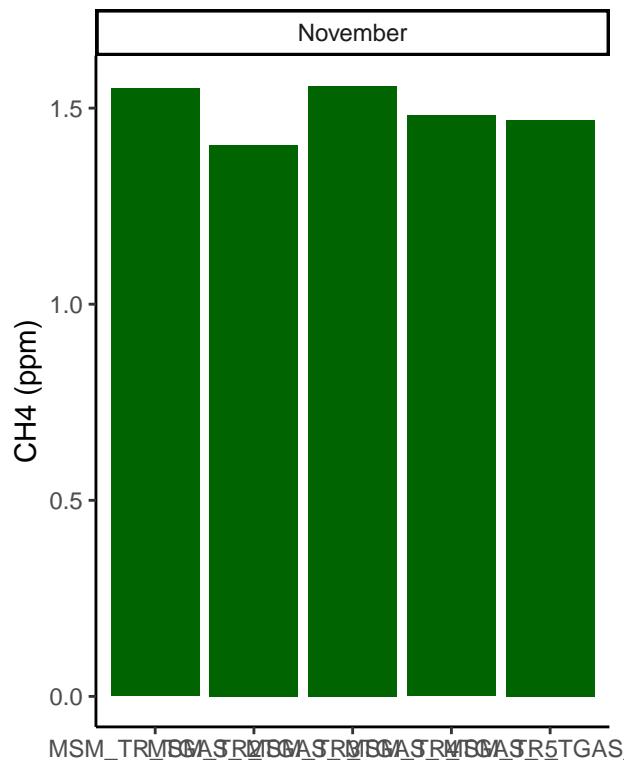
#check results
#head(Samples)

#quick first look at the samples
ch4_samples <- ggplot(data = Samples, aes(x = Sample_ID, y = CH4_Conc_ppm, fill=CH4_Flag)) +
  geom_bar(stat = 'identity') +
  scale_fill_manual(values=c( "darkgreen", "red"))+
  #scale_fill_gradient2(low='red', mid='white', high='blue', space='Lab') +
  theme_classic() + labs(x= " ", y="CH4 (ppm)", title="CH4: Green = Within Range") +
  theme(legend.position="none") +
  facet_grid(~Sample_Month)

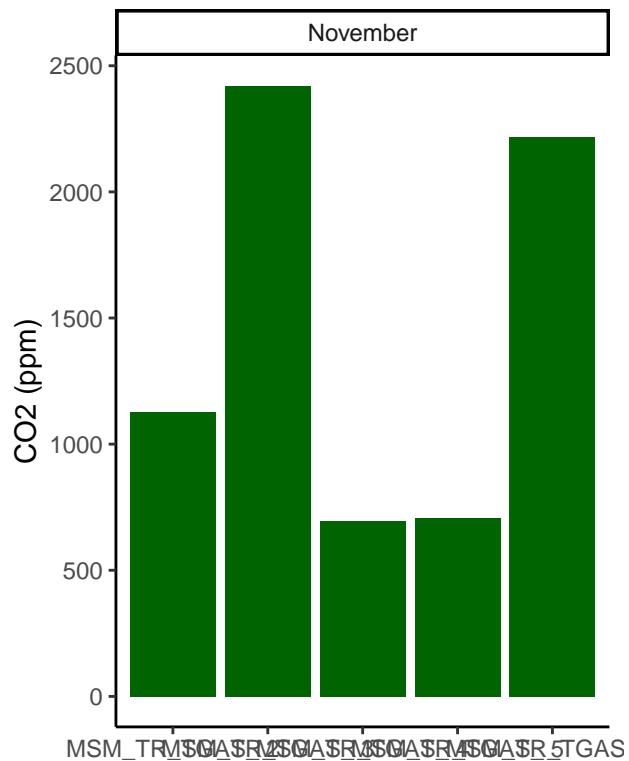
co2_samples <- ggplot(data = Samples, aes(x = Sample_ID, y = CO2_Conc_ppm, fill=CO2_Flag)) +
  geom_bar(stat = 'identity') +
  scale_fill_manual(values=c("darkgreen", "red"))+
  #scale_fill_gradient2(low='red', mid='white', high='blue', space='Lab') +
  theme_classic() + labs(x= " ", y="CO2 (ppm)", title="CO2: Green = Within Range") +
  theme(legend.position="none") +
  facet_grid(~Sample_Month)

ggarrange(ch4_samples, co2_samples, nrow=1, ncol=2)
```

CH4: Green = Within Range



CO2: Green = Within Range



Write out processed data & slopes

```
#check results
head(Samples)
```

```
##      Machine User Run_Date Sample_Type Type1 Sample_Year Sample_Month
## 1 Varian GC Wegner 20240311 Unknown    TGW      2023      November
## 2 Varian GC Wegner 20240311 Unknown    TGW      2023      November
## 3 Varian GC Wegner 20240311 Unknown    TGW      2023      November
## 4 Varian GC Wegner 20240311 Unknown    TGW      2023      November
## 5 Varian GC Wegner 20240311 Unknown    TGW      2023      November
##           Sample_ID Dilution_Factor STD_Conc CO2_Area CH4_Area Field.Notes
## 1 MSM_TR_TGAS_ 2          2.83     NA 178484     868       NA
## 2 MSM_TR_TGAS_ 3          2.83     NA 374692     772       NA
## 3 MSM_TR_TGAS_ 4          2.83     NA 112466     872       NA
## 4 MSM_TR_TGAS_ 5          2.83     NA 114307     822       NA
## 5 MSM_TR_TGAS_ 6          2.83     NA 344017     815       NA
##                                         Lab.Notes CH4_Curve
## 1 these got skipped on the first run; 10mL N2 added in the lab      Low
## 2                                         10mL N2 added in the lab      Low
## 3                                         10mL N2 added in the lab      Low
## 4                                         10mL N2 added in the lab      Low
## 5                                         10mL N2 added in the lab      Low
##      CH4_Conc_ppm CO2_Conc_ppm     CH4_Flag     CO2_Flag CH4_Conc_ppm_dilcorr
```

```

## 1      1.550096 1128.6891 Within Range Within Range          4.386772
## 2      1.406194 2419.1595 Within Range Within Range          3.979529
## 3      1.556092 694.4853 Within Range Within Range          4.403740
## 4      1.481143 706.5936 Within Range Within Range          4.191634
## 5      1.470650 2217.4084 Within Range Within Range          4.161940
##   CO2_Conc_ppm_dilcorr
## 1            3194.190
## 2            6846.221
## 3            1965.393
## 4            1999.660
## 5            6275.266

```

```

#pull out what we need
Samples1 <- Samples[ ,c(1:3,5:9,13, 18:21)]
head(Samples1)

```

```

##   Machine User Run_Date Type1 Sample_Year Sample_Month       Sample_ID
## 1 Varian GC Wegner 20240311    TGW      2023 November MSM_TR_TGAS_ 2
## 2 Varian GC Wegner 20240311    TGW      2023 November MSM_TR_TGAS_ 3
## 3 Varian GC Wegner 20240311    TGW      2023 November MSM_TR_TGAS_ 4
## 4 Varian GC Wegner 20240311    TGW      2023 November MSM_TR_TGAS_ 5
## 5 Varian GC Wegner 20240311    TGW      2023 November MSM_TR_TGAS_ 6
##   Dilution_Factor Field.Notes     CH4_Flag     CO2_Flag CH4_Conc_ppm_dilcorr
## 1            2.83        NA Within Range Within Range          4.386772
## 2            2.83        NA Within Range Within Range          3.979529
## 3            2.83        NA Within Range Within Range          4.403740
## 4            2.83        NA Within Range Within Range          4.191634
## 5            2.83        NA Within Range Within Range          4.161940
##   CO2_Conc_ppm_dilcorr
## 1            3194.190
## 2            6846.221
## 3            1965.393
## 4            1999.660
## 5            6275.266

```

```

Samples1 <- Samples1 %>%
  separate(Sample_ID, into = c("Site", "Gas_Sample", "Zone", "Replicate"), sep = "_", remove = FALSE) %>%
  mutate(Tree_Code = "SF") %>%
  mutate(Tree_Info = case_when(
    Tree_Code == "DS" ~ "Dead Standing",
    Tree_Code == "SF" ~ "Sapflow Monitoring",
    TRUE ~ "Other" # Optional: handles any values that aren't DS or SF
  )) %>%
  mutate(Status = case_when(
    Tree_Code == "DS" ~ "Dead Standing",
    Tree_Code == "SF" ~ "Living",
    TRUE ~ "Other"
  )) %>%
  mutate(Project = "COMPASS: Synoptic",
         Region = "CB") %>%
  rename( Year = Sample_Year,
         Month = Sample_Month,
         CH4_ppm = CH4_Conc_ppm_dilcorr ,

```

```

CO2_ppm = CO2_Conc_ppm_dilcorr ) %>%
mutate(CH4_Flag = case_when(
  CH4_Flag == "Needs_Dilution" ~ "Over Std Curve Range",
  TRUE ~ "Within Std Curve Range"
)) %>%
mutate(CO2_Flag = case_when(
  CO2_Flag == "Needs_Dilution" ~ "Over Std Curve Range",
  TRUE ~ "Within Std Curve Range"
))

final <- Samples1 %>%
  select( "Project", "Region" , "Year", "Month" , "Site", "Zone", "Gas_Sample",
  "Sample_ID", "Tree_Code", "Replicate", "Status", "Tree_Info",
  "CH4_ppm", "CH4_Flag", "CO2_ppm", "CO2_Flag")

write.csv(final, "Processed Data/COMPASS_Synoptic_TGW_202311_Processed.csv")

```

#end