

COMPASS_Synoptic_TGW_2022: June

Stephanie J. Wilson

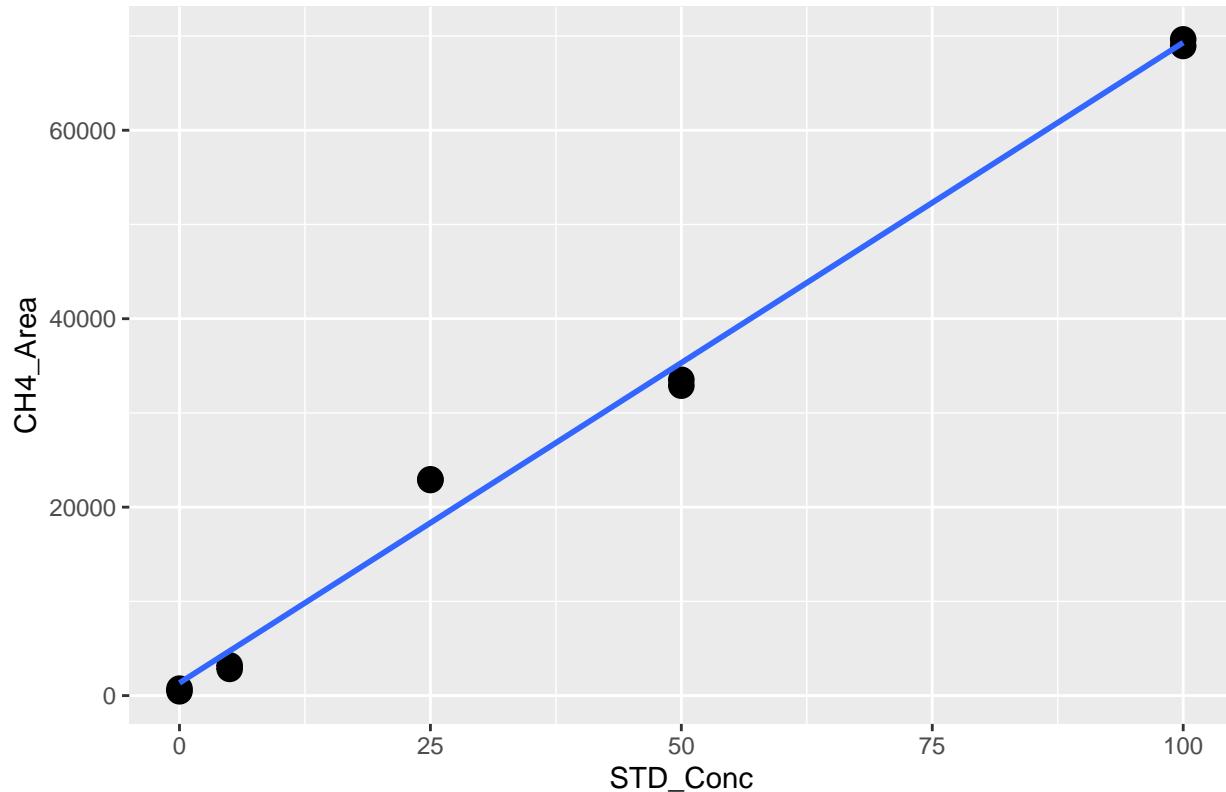
2023-01-14

##Set Up

Read in first data file and assess standard curves

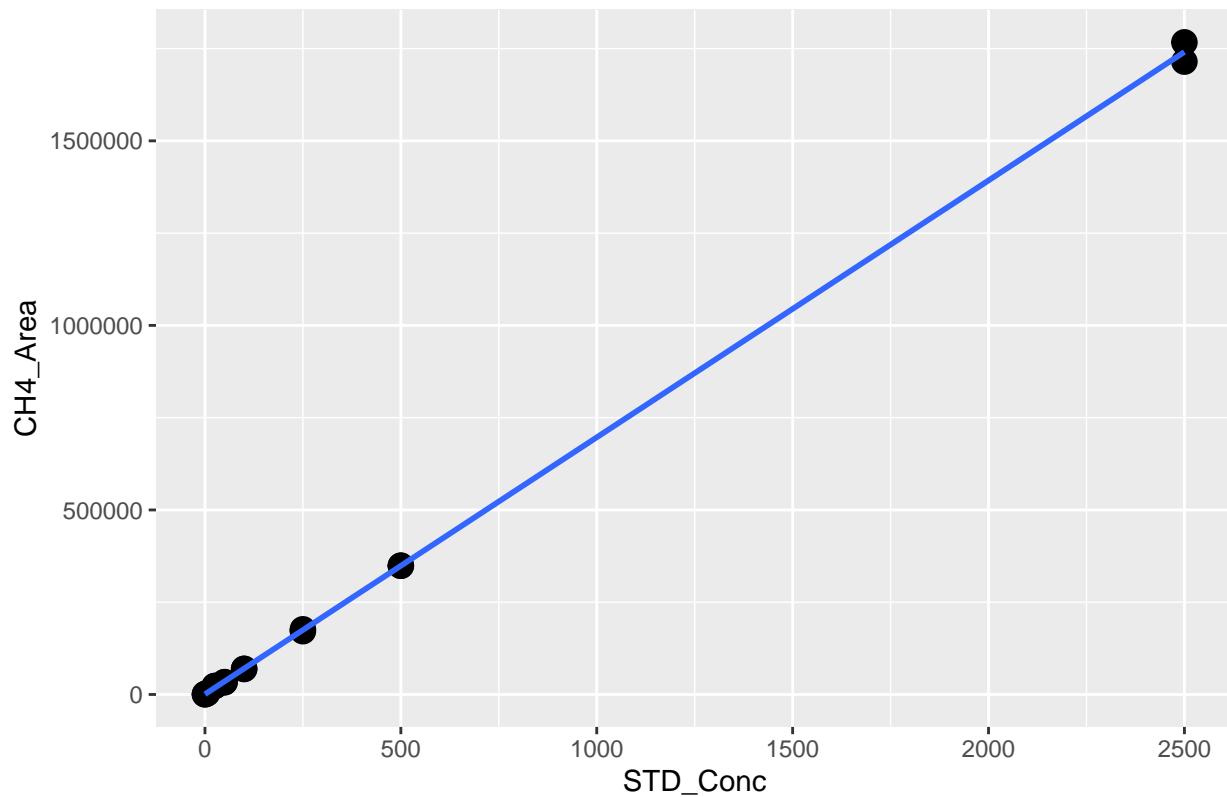
```
##      Machine          User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202206     Blank   TGAS      2022
## 2 Varian GC Stephanie J. Wilson 202206 Unknown   TGAS      2022
## 3 Varian GC Stephanie J. Wilson 202206 Unknown   TGAS      2022
## 4 Varian GC Stephanie J. Wilson 202206 Unknown   TGAS      2022
## 5 Varian GC Stephanie J. Wilson 202206 Unknown   TGAS      2022
## 6 Varian GC Stephanie J. Wilson 202206 Unknown   TGAS      2022
##      Sample_Month    Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1       June        Blank_1             1       NA      0      0
## 2       June  MSM_TGAS_UP_SF_1         2       NA  12697 1791814
## 3       June  MSM_TGAS_UP_SF_2         2       NA  23232 5697789
## 4       June  MSM_TGAS_UP_SF_3         2       NA   8442 10718524
## 5       June  MSM_TGAS_UP_SF_4         2       NA   8731 6094426
## 6       June  MSM_TGAS_UP_SF_5         2       NA  34611 8661859
##      Field.Notes Lab.Notes
## 1       NA        NA
## 2       NA        NA
## 3       NA        NA
## 4       NA        NA
## 5       NA        NA
## 6       NA        NA
## `geom_smooth()` using formula = 'y ~ x'
```

CH4 LOW Std Curve



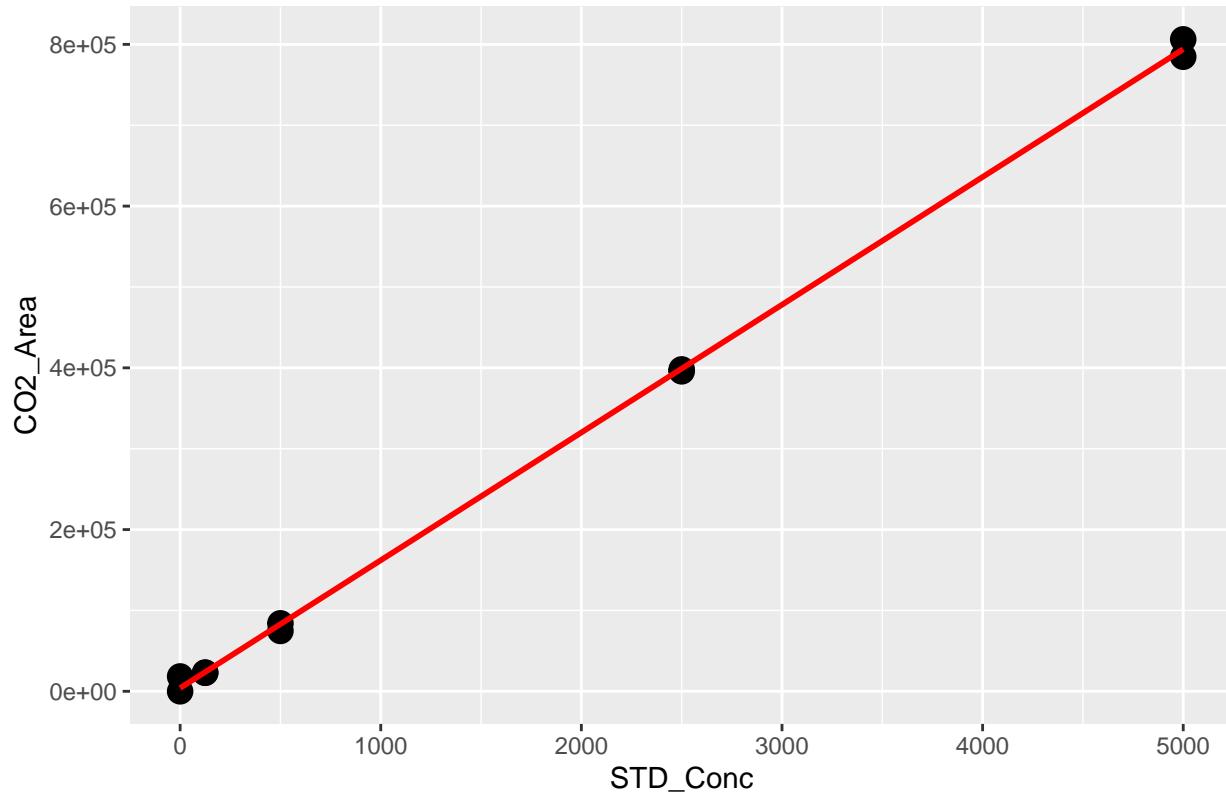
```
##  
## Call:  
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -2430.8 -1733.7 - 744.1   173.6  4623.8  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (Intercept)           1342.56     1201.65   1.117   0.296  
## stds_ch4_low$STD_Conc  679.55      23.43  29.001 2.16e-09 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 2706 on 8 degrees of freedom  
## Multiple R-squared:  0.9906, Adjusted R-squared:  0.9894  
## F-statistic: 841.1 on 1 and 8 DF,  p-value: 2.163e-09  
  
## 'geom_smooth()' using formula = 'y ~ x'
```

CH4 HIGH Std Curve



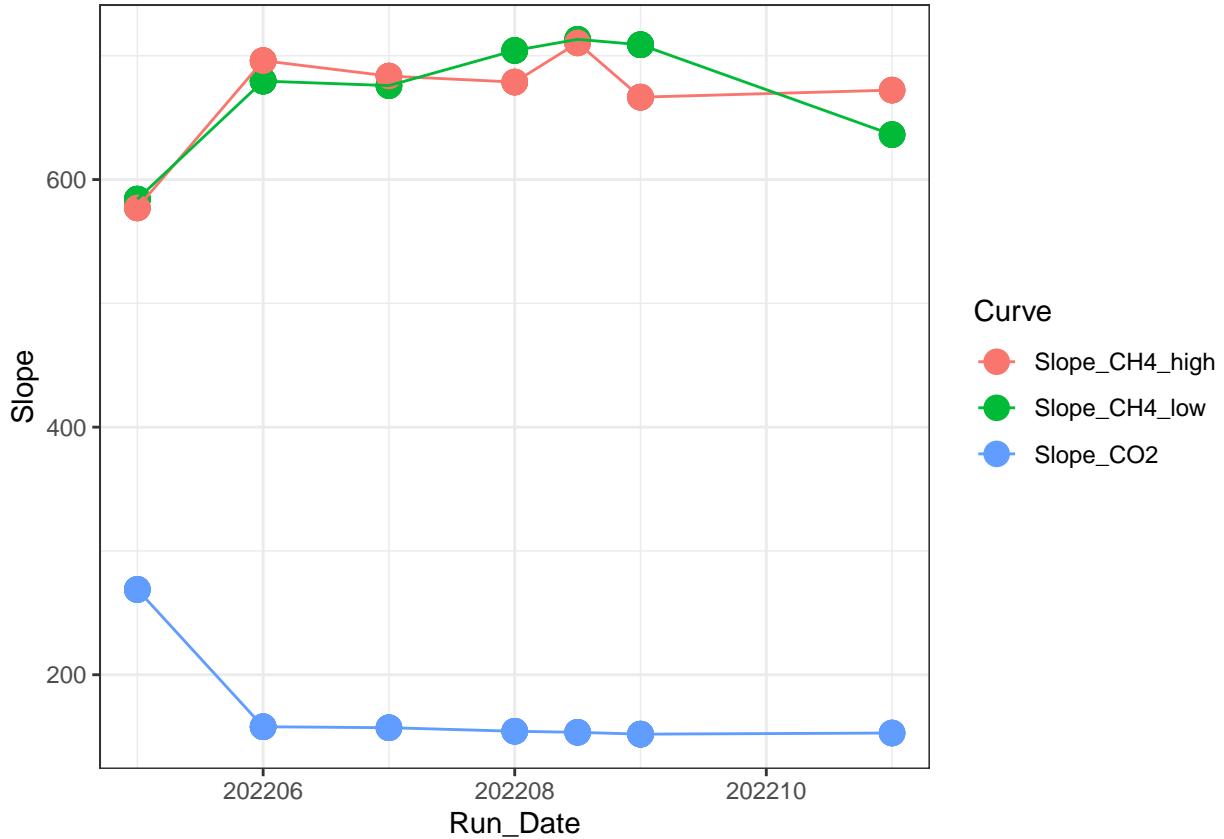
```
##  
## Call:  
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)  
##  
## Residuals:  
##      Min       1Q     Median       3Q      Max  
## -26065.1  -1479.7   -414.3   670.4  26209.9  
##  
## Coefficients:  
##                         Estimate Std. Error t value Pr(>|t|)  
## (Intercept)      664.824    2880.168   0.231   0.821  
## stds_ch4$STD_Conc 696.019      3.177 219.093 <2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 10150 on 14 degrees of freedom  
## Multiple R-squared:  0.9997, Adjusted R-squared:  0.9997  
## F-statistic: 4.8e+04 on 1 and 14 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:
##      Min       1Q   Median       3Q      Max 
## -9295.7 -3873.2  -928.8   611.6 14324.4 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 4049.58    3415.57   1.186   0.27    
## stds_co2$STD_Conc 157.99      1.36 116.130 3.38e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8233 on 8 degrees of freedom
## Multiple R-squared:  0.9994, Adjusted R-squared:  0.9993 
## F-statistic: 1.349e+04 on 1 and 8 DF,  p-value: 3.379e-14

##      X          Curve      R2      Slope Intercept Run_Date
## 1 1 Slope_CH4_low 0.9989050 584.1700  292.6667 202205
## 2 2 Slope_CH4_high 0.9988404 577.1458 -1829.3469 202205
## 3 3 Slope_CO2 0.8941001 268.8803 82323.2843 202205
## 4 4 Slope_CH4_low 0.9989050 584.1700  292.6667 202205
## 5 5 Slope_CH4_high 0.9988404 577.1458 -1829.3469 202205
## 6 6 Slope_CO2 0.8941001 268.8803 82323.2843 202205
```



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_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)

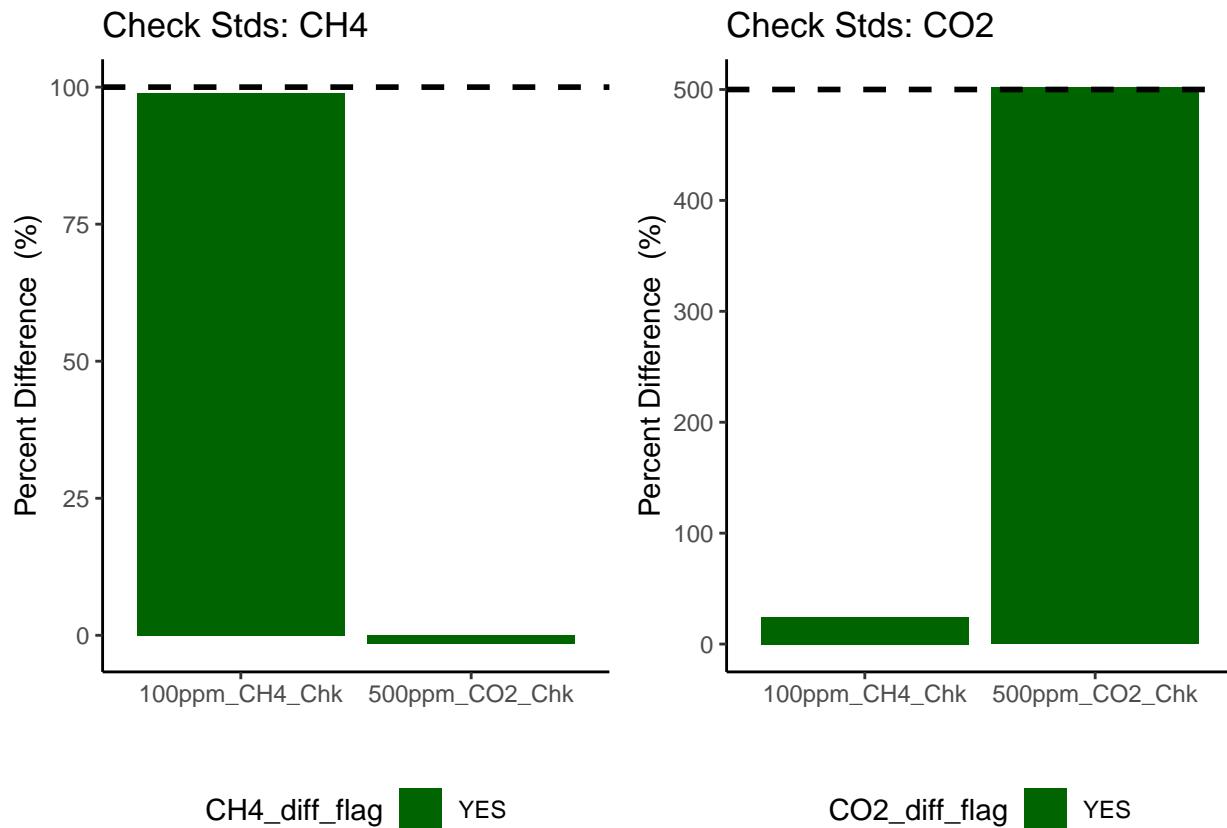
```

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)

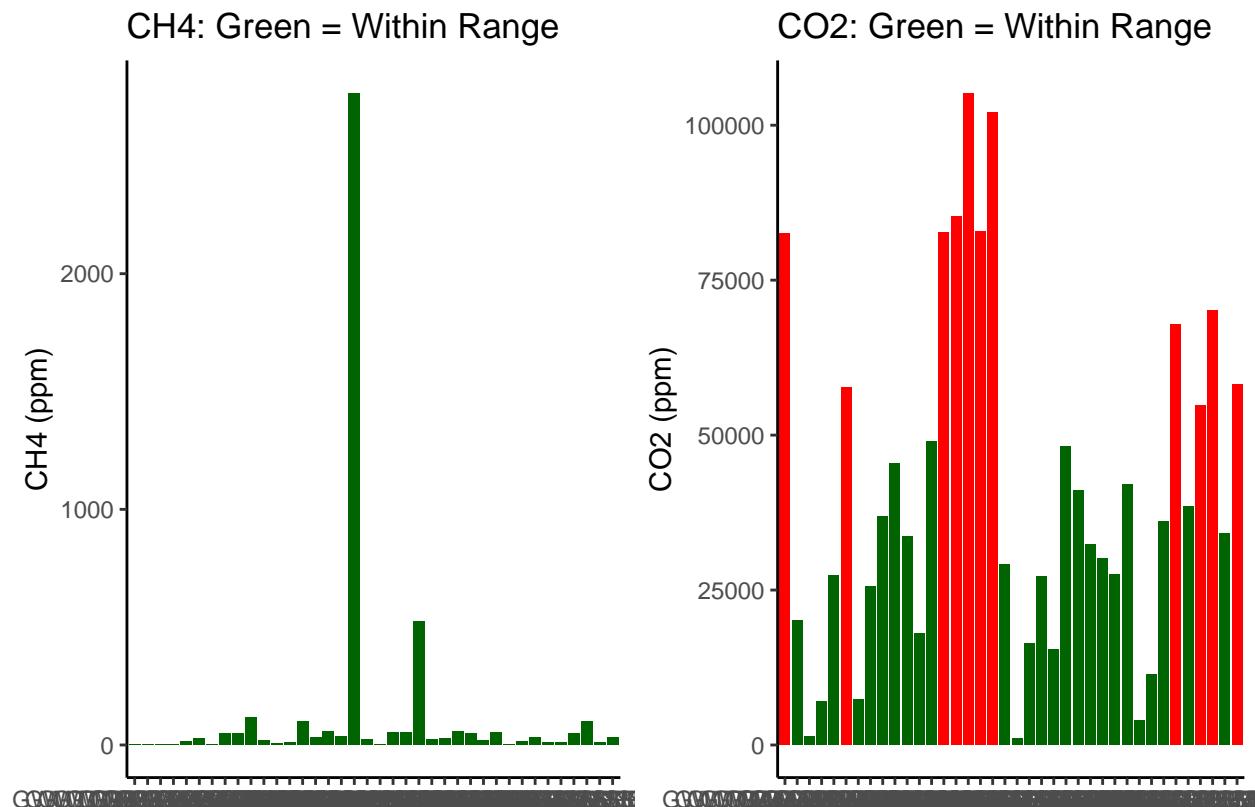
#check results
#head(Samples)

#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")

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( "red","darkgreen"))+
  #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")

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

```



Write out processed data & slopes

```
#check results
head(Samples)

##      Machine          User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
## 2 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
## 3 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
## 4 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
## 5 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
## 6 Varian GC Stephanie J. Wilson 202206 Unknown    TGAS     2022
##   Sample_Month       Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1       June MSM_TGAS_UP_SF_1                  2      NA 12697 1791814
## 2       June MSM_TGAS_UP_SF_2                  2      NA 23232 5697789
## 3       June MSM_TGAS_UP_SF_3                  2      NA 8442 10718524
## 4       June MSM_TGAS_UP_SF_4                  2      NA 8731 6094426
## 5       June MSM_TGAS_UP_SF_5                  2      NA 34611 8661859
## 6       June MSM_TGAS_UP_SF_6                  2      NA 68050 11080250
##   Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm      CH4_Flag
## 1        NA        NA      Low    16.70887 11315.66 Within Range
## 2        NA        NA      Low    32.21188 36038.55 Within Range
## 3        NA        NA      Low    10.44734 67817.32 Within Range
## 4        NA        NA      Low    10.87262 38549.07 Within Range
## 5        NA        NA      Low    48.95689 54799.65 Within Range
## 6        NA        NA      Low    98.16478 70106.87 Within Range
##           CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range            33.41775          22631.33
## 2 Within Range            64.42376          72077.11
## 3 Needs Dilution        20.89467          135634.65
## 4 Within Range            21.74524          77098.14
## 5 Needs Dilution         97.91379          109599.30
## 6 Needs Dilution        196.32956          140213.74
```

```
#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
## 1 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
## 2 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
## 3 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
## 4 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
## 5 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
## 6 Varian GC Stephanie J. Wilson 202206    TGAS     2022       June
##   Sample_ID Dilution_Factor Field.Notes      CH4_Flag      CO2_Flag
## 1 MSM_TGAS_UP_SF_1                  2      NA Within Range Within Range
## 2 MSM_TGAS_UP_SF_2                  2      NA Within Range Within Range
## 3 MSM_TGAS_UP_SF_3                  2      NA Within Range Needs Dilution
## 4 MSM_TGAS_UP_SF_4                  2      NA Within Range Within Range
## 5 MSM_TGAS_UP_SF_5                  2      NA Within Range Needs Dilution
## 6 MSM_TGAS_UP_SF_6                  2      NA Within Range Needs Dilution
```

```

##    CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1            33.41775      22631.33
## 2            64.42376      72077.11
## 3            20.89467      135634.65
## 4            21.74524      77098.14
## 5            97.91379     109599.30
## 6           196.32956     140213.74

Samples1 <- Samples1 %>%
  separate(Sample_ID, into = c("Site", "Gas_Sample", "Zone", "Tree_Code", "Replicate"), sep = "_", remove = TRUE) %>%
  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_202206_Processed.csv")

#end

```