

# COMPASS\_Synoptic\_TGW\_2022: Sept & Oct part 1

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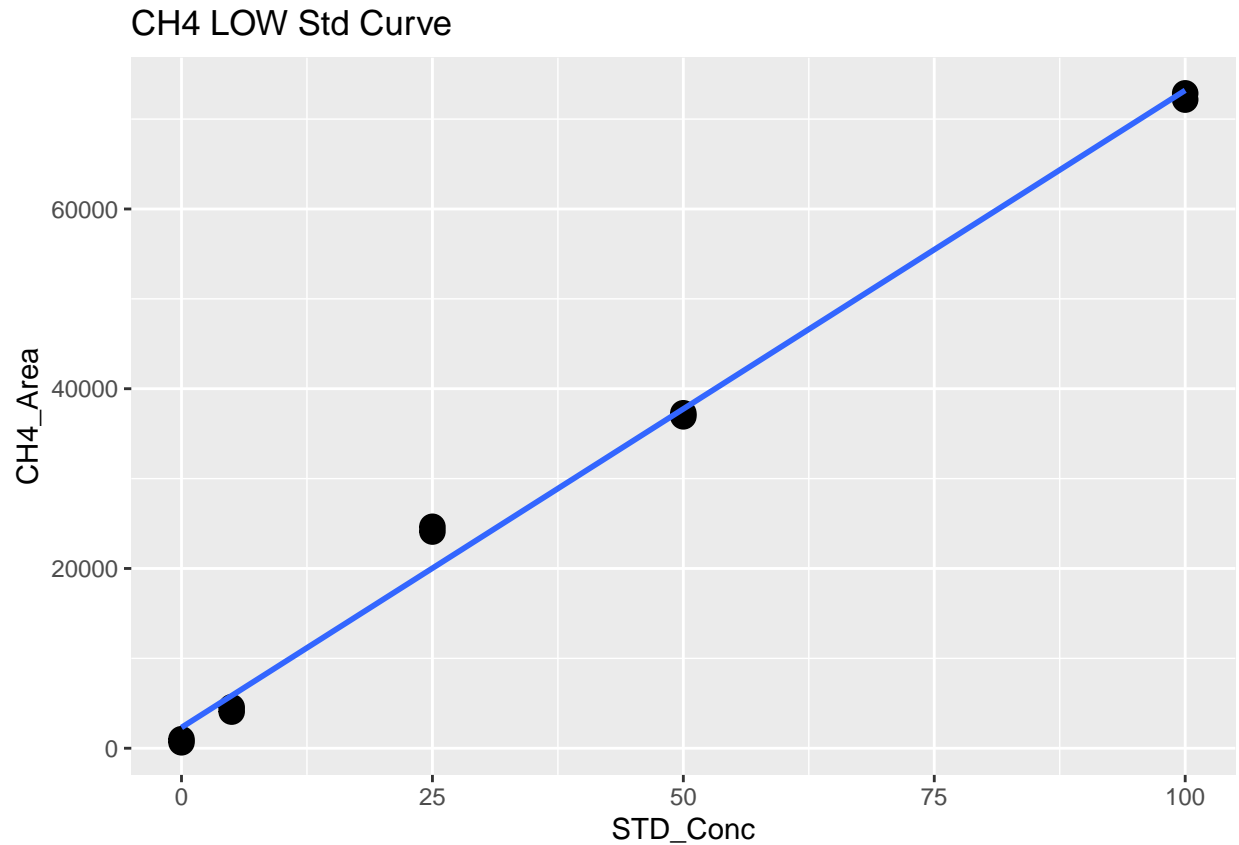
2023-01-14

##Set Up

Read in first data file and assess standard curves

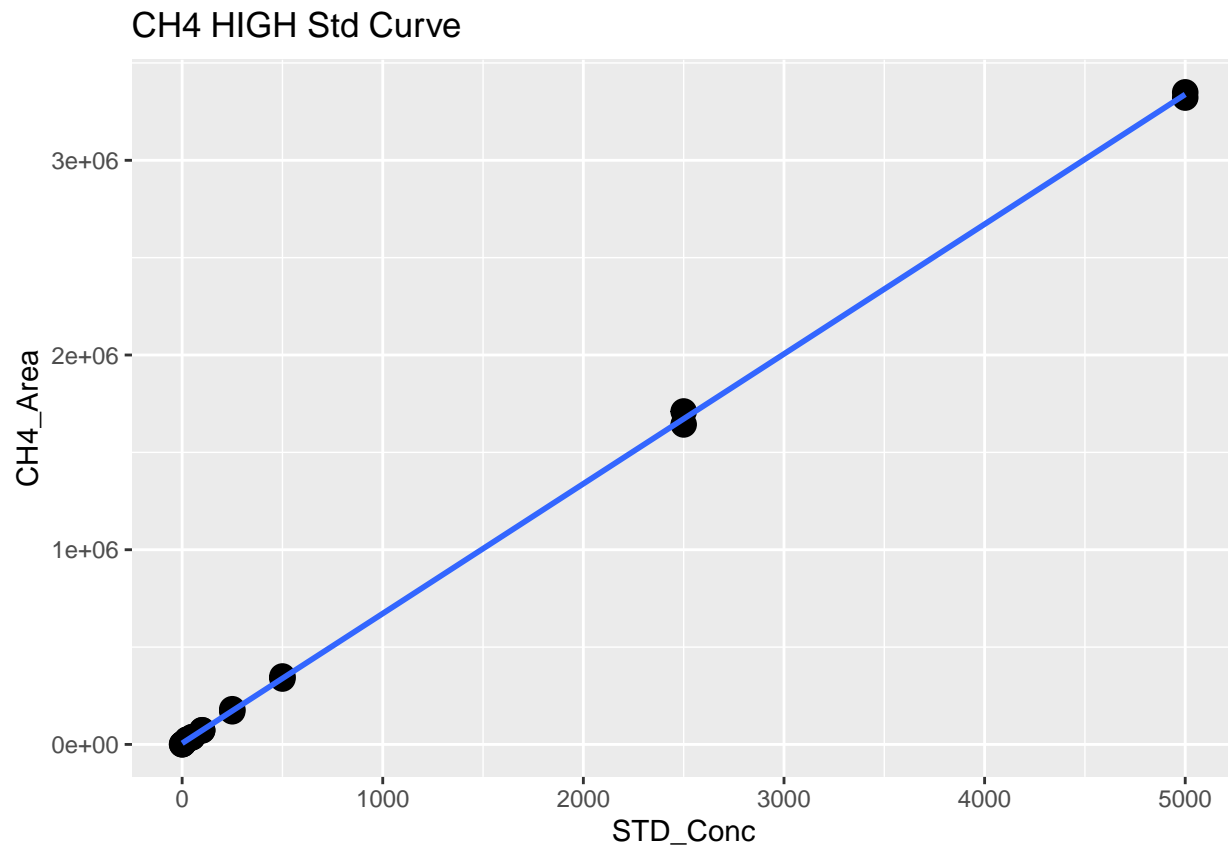
```
##      Machine      User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202212 Chk_STD TGAS 2022
## 2 Varian GC Stephanie J. Wilson 202212 Chk_STD TGAS 2022
## 3 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 4 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 5 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 6 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      <NA>      100ppm chk      1      100      61383      NA
## 2      <NA>      500ppm CO2 chk      1      500      NA      93076
## 3      September MSM_TGAS_UP_SF_1      2      NA      31673 4381503
## 4      September MSM_TGAS_UP_SF_2      2      NA      66560 4874844
## 5      September MSM_TGAS_UP_SF_3      2      NA      11339 7003458
## 6      September MSM_TGAS_UP_SF_4      2      NA      9099 5176853
##      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'
```



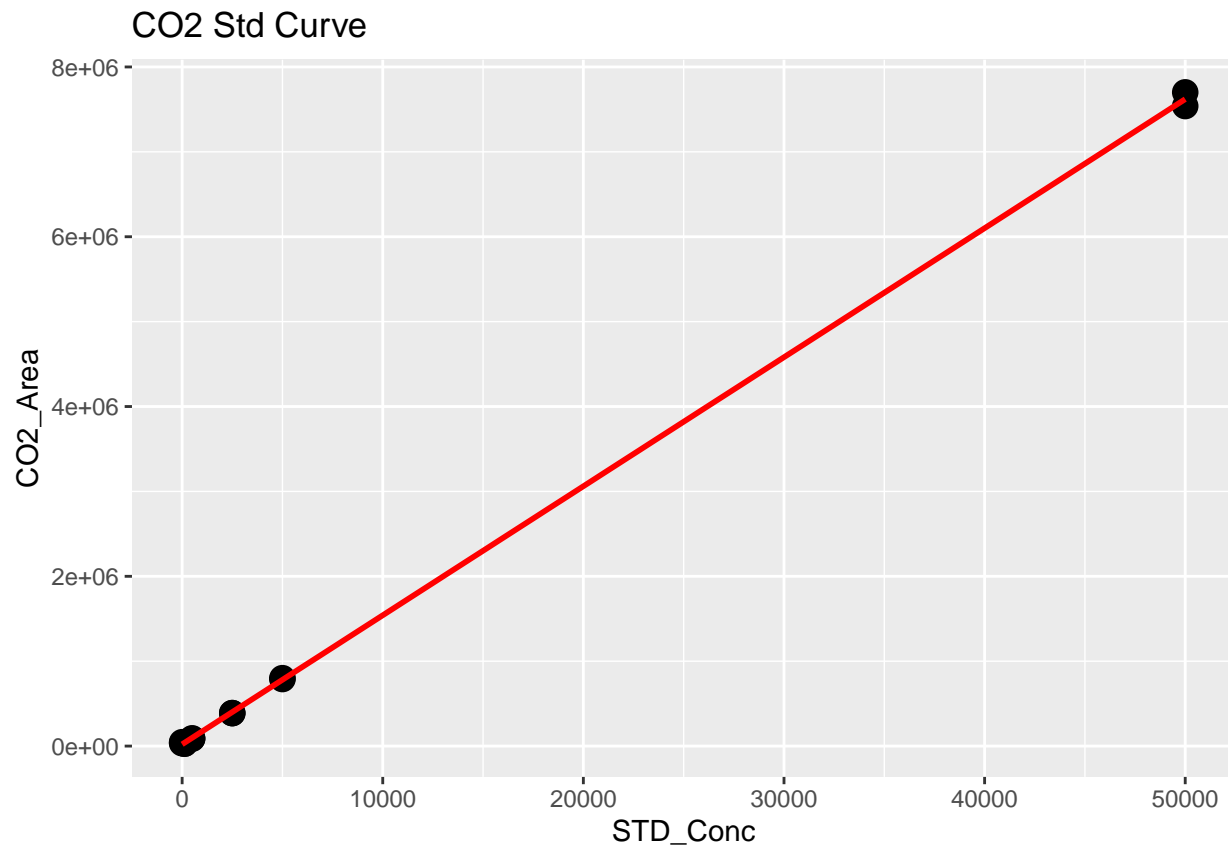
```
##
## Call:
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -1777.5 -1300.3  -928.8  -370.2   4611.0
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2306.85    1101.37     2.095  0.0695 .
## stds_ch4_low$STD_Conc    708.93      21.48    33.010 7.74e-10 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2480 on 8 degrees of freedom
## Multiple R-squared:  0.9927, Adjusted R-squared:  0.9918
## F-statistic: 1090 on 1 and 8 DF, p-value: 7.738e-10

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



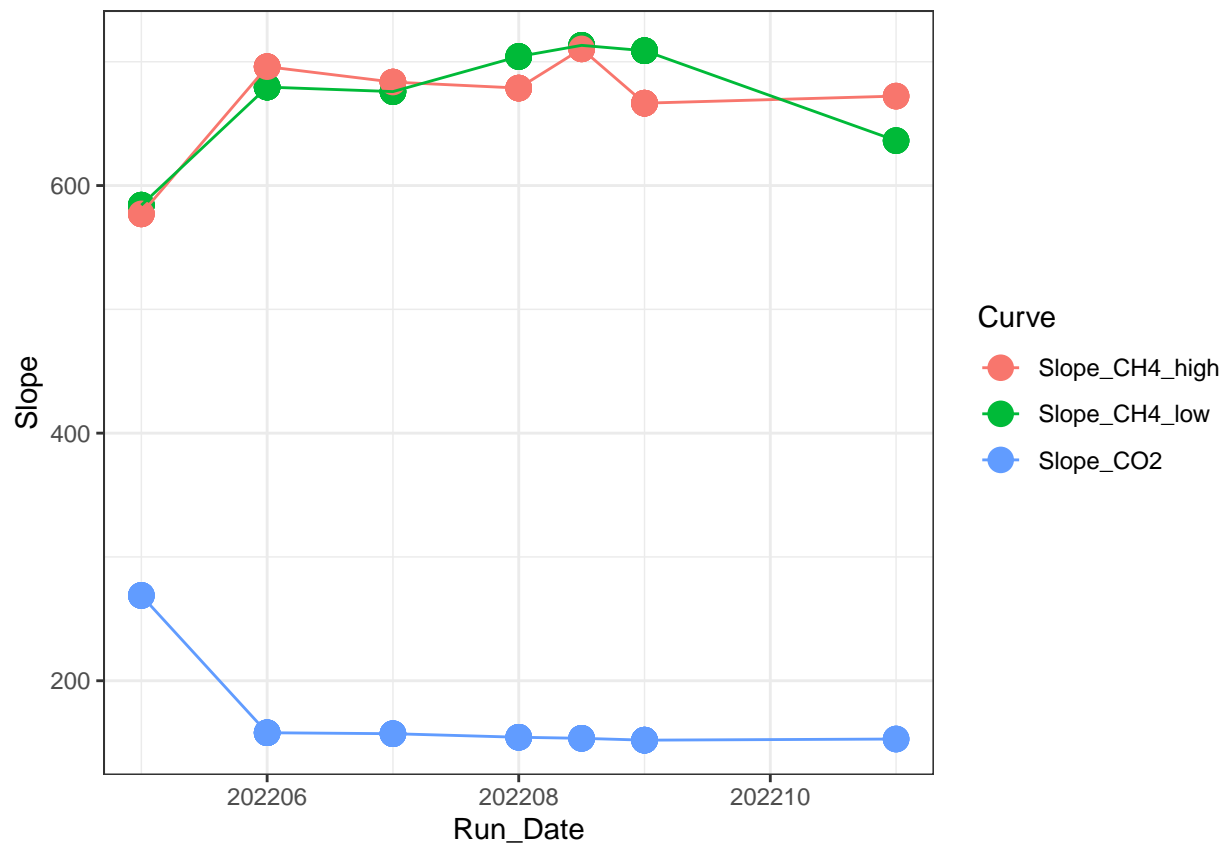
```
##
## Call:
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -28677  -4679  -1195    2106   37909
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    5738.283    3679.618     1.559   0.138
## stds_ch4$STD_Conc  666.556      1.964  339.304 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 13520 on 16 degrees of freedom
## Multiple R-squared:  0.9999, Adjusted R-squared:  0.9999
## F-statistic: 1.151e+05 on 1 and 16 DF, p-value: < 2.2e-16

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



```
##
## Call:
## lm(formula = stds_co2$CO2_Area ~ stds_co2$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -81507 -11491  -8260   14038   80240
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2.104e+04  1.267e+04    1.66   0.128
## stds_co2$STD_Conc 1.520e+02  6.169e-01  246.38 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 38710 on 10 degrees of freedom
## Multiple R-squared:  0.9998, Adjusted R-squared:  0.9998
## F-statistic: 6.07e+04 on 1 and 10 DF,  p-value: < 2.2e-16

##      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)

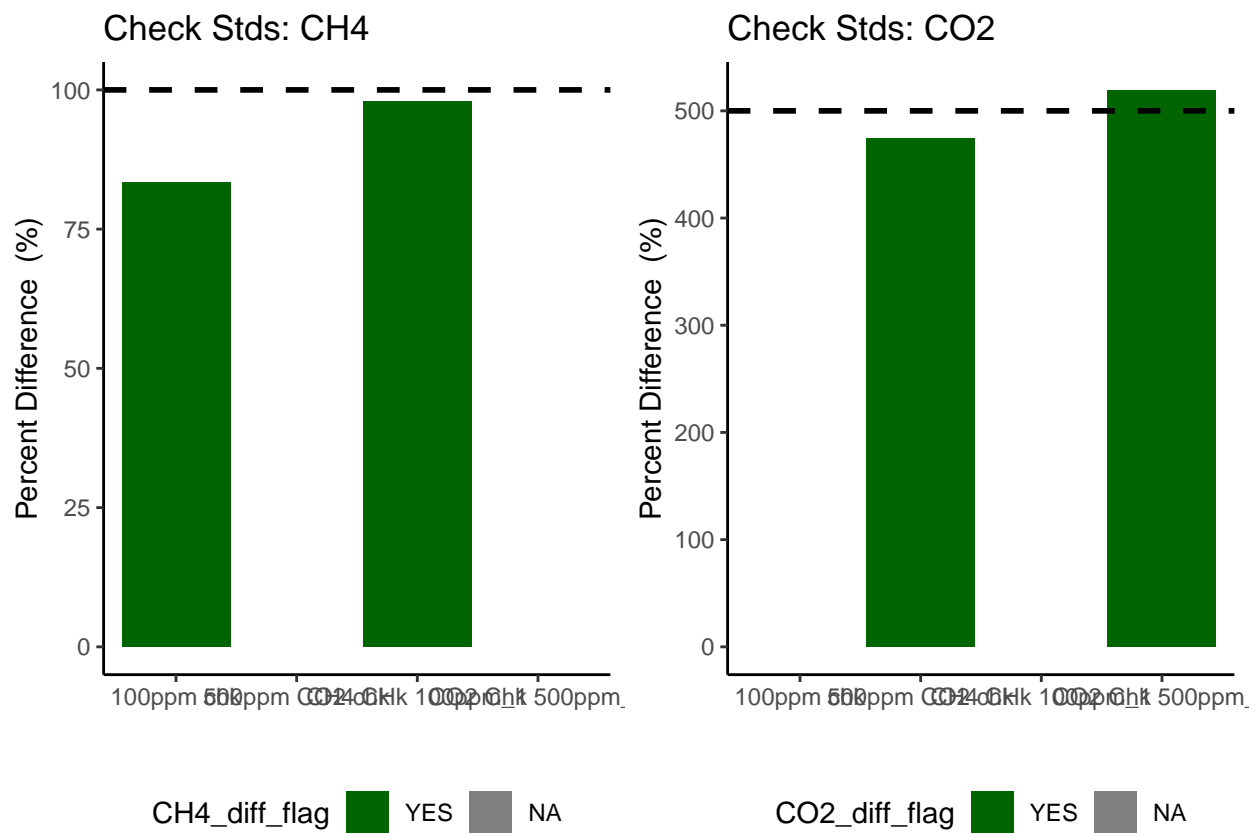
#####make flags for any dilutions needed
#highest CH4 standard = 10000
#highest CO2 standard = 50000

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.

## Warning: Removed 2 rows containing missing values or values outside the scale range
## ('geom_bar()').
## Removed 2 rows containing missing values or values outside the scale range
## ('geom_bar()').
```



## Dilution correct samples

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

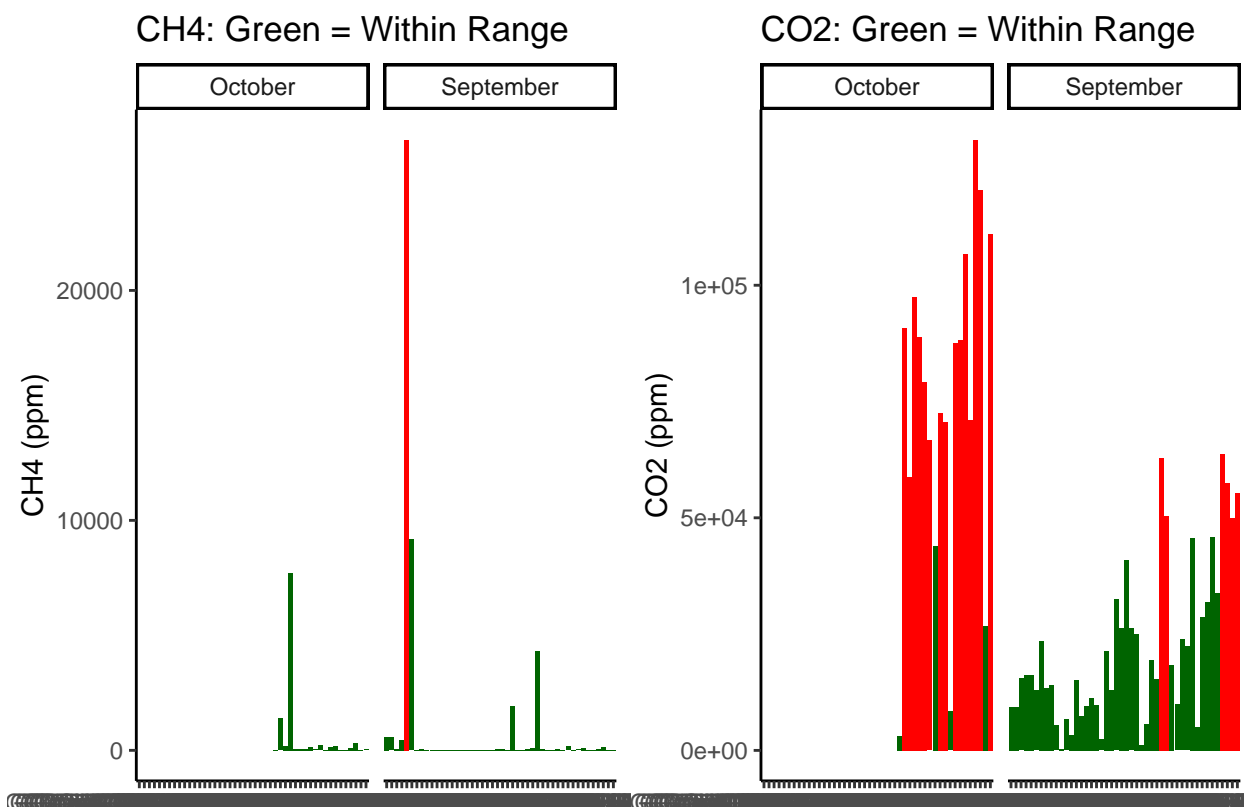
Samples$CO2_Conc_ppm_dilcorr <- (Samples$CO2_Conc_ppm * Samples$Dilution_Factor)

#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( "red","darkgreen" ))+
  #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("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") +
  facet_grid(~Sample_Month)

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 202212 Unknown TGAS 2022
## 2 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 3 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 4 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 5 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## 6 Varian GC Stephanie J. Wilson 202212 Unknown TGAS 2022
## Sample_Month Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1 September MSM_TGAS_UP_SF_1 2 NA 31673 4381503
## 2 September MSM_TGAS_UP_SF_2 2 NA 66560 4874844
## 3 September MSM_TGAS_UP_SF_3 2 NA 11339 7003458
## 4 September MSM_TGAS_UP_SF_4 2 NA 9099 5176853
## 5 September MSM_TGAS_UP_SF_5 2 NA 37421 9708735
## 6 September MSM_TGAS_UP_SF_6 2 NA 93715 8766135
## Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm CH4_Flag
## 1 NA NA Low 41.423412 28688.29 Within Range
## 2 NA NA Low 90.634445 31934.07 Within Range
## 3 NA NA Low 12.740603 45938.61 Within Range
## 4 NA NA Low 9.580896 33921.04 Within Range
```



```
## 5      NA      NA      Low      49.531447      63737.13 Within Range
## 6      NA      NA      High     131.987083      57535.59 Within Range
##      CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range      82.84682      57376.58
## 2 Within Range     181.26889      63868.14
## 3 Within Range     25.48121      91877.23
## 4 Within Range     19.16179      67842.09
## 5 Needs Dilution     99.06289     127474.26
## 6 Needs Dilution    263.97417     115071.18
```

```
#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 202212 TGAS      2022      September
## 2 Varian GC Stephanie J. Wilson 202212 TGAS      2022      September
## 3 Varian GC Stephanie J. Wilson 202212 TGAS      2022      September
## 4 Varian GC Stephanie J. Wilson 202212 TGAS      2022      September
## 5 Varian GC Stephanie J. Wilson 202212 TGAS      2022      September
## 6 Varian GC Stephanie J. Wilson 202212 TGAS      2022      September
##      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 Within Range
## 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      82.84682      57376.58
## 2     181.26889      63868.14
## 3     25.48121      91877.23
## 4     19.16179      67842.09
## 5     99.06289     127474.26
## 6    263.97417     115071.18
```

```
Samples1 <- Samples1 %>%
  separate(Sample_ID, into = c("Site", "Gas_Sample", "Zone", "Tree_Code", "Replicate"), sep = "_", remove = FALSE)
  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_202209-10_Processed.csv")

```

#end