

# COMPASS\_Synoptic\_TGW\_2022: Oct part 2 & Nov

Stephanie J. Wilson

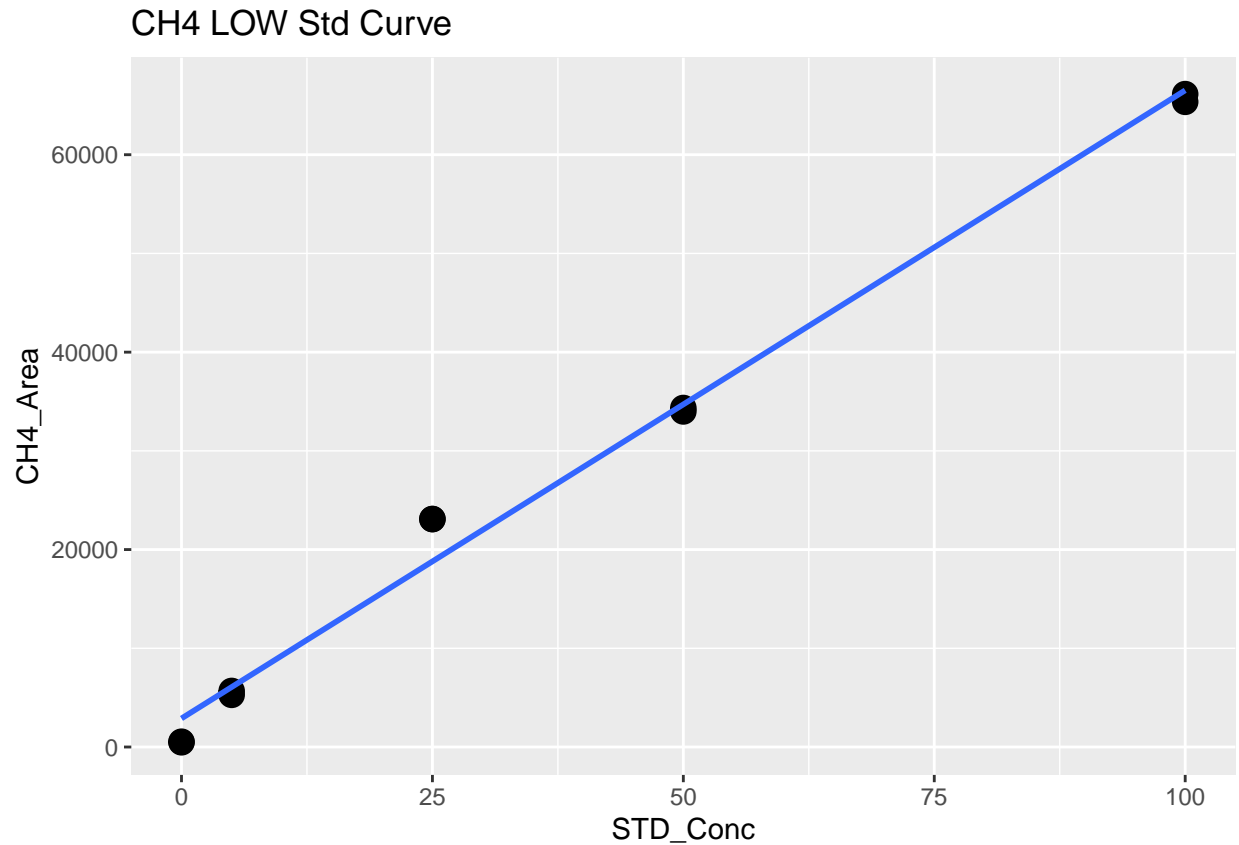
2023-01-14

##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 202212.5      Blank  TGAS      2022
## 2 Varian GC Stephanie J. Wilson 202212.5      Chk_STD TGAS      2022
## 3 Varian GC Stephanie J. Wilson 202212.5      Chk_STD TGAS      2022
## 4 Varian GC Stephanie J. Wilson 202212.5      Unknown TGAS      2022
## 5 Varian GC Stephanie J. Wilson 202212.5      Unknown TGAS      2022
## 6 Varian GC Stephanie J. Wilson 202212.5      Unknown TGAS      2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      <NA>      Blank      1      NA      391      18821
## 2      <NA>      CH4 Chk 100ppm      1      100      65160      NA
## 3      <NA>      CO2 Chk 500ppm      1      500      NA      86005
## 4      October GWI_TGAS_UP_SF_1      2      NA      18193 3989643
## 5      October GWI_TGAS_UP_SF_2      2      NA      22122 6882701
## 6      October GWI_TGAS_UP_SF_3      2      NA      21739 5279576
##      Field.Notes Lab.Notes
## 1      NA      NA
## 2      NA      NA
## 3      NA      NA
## 4      NA      NA
## 5      NA      NA
## 6      NA      NA

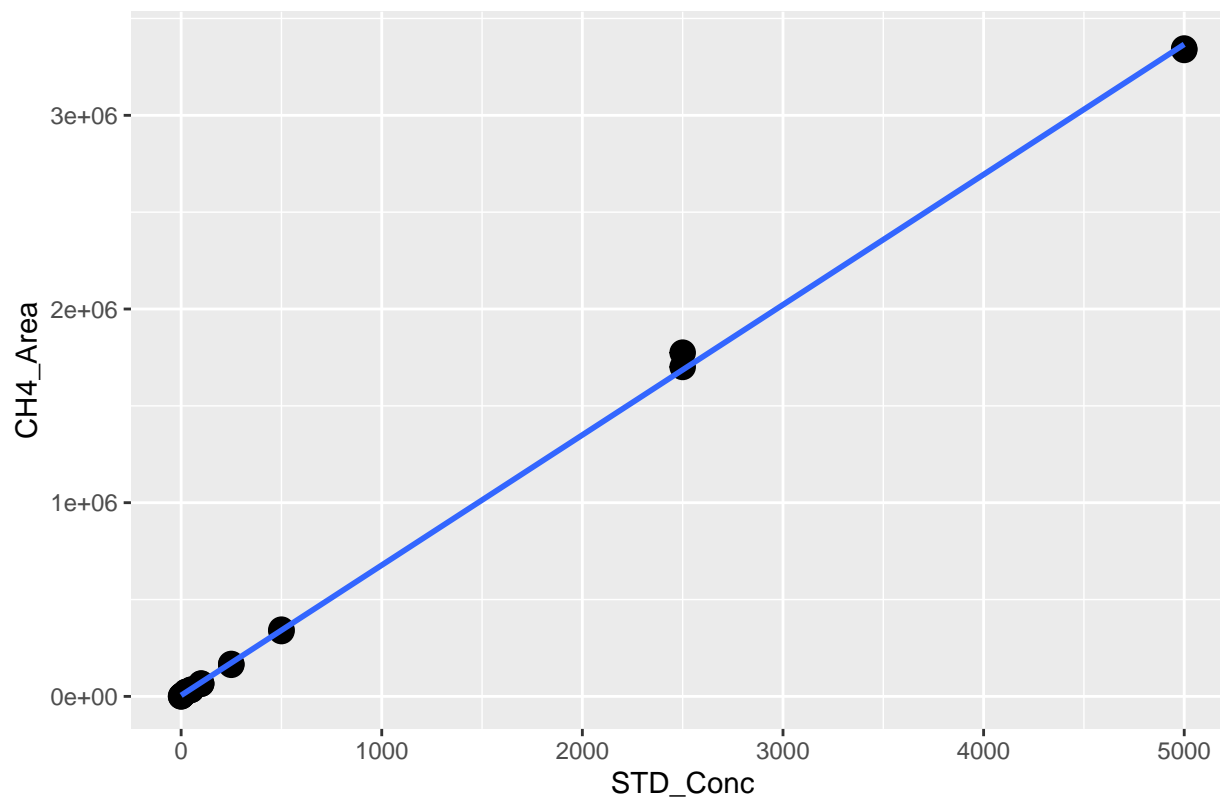
## 'geom_smooth()' using formula = 'y ~ x'
```



```
##
## Call:
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2416.3 -1073.6  -554.3   -367.8   4298.4
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    2893.27    1122.74   2.577  0.0328 *
## stds_ch4_low$STD_Conc  636.29      21.89  29.064 2.13e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2529 on 8 degrees of freedom
## Multiple R-squared:  0.9906, Adjusted R-squared:  0.9894
## F-statistic: 844.7 on 1 and 8 DF, p-value: 2.126e-09

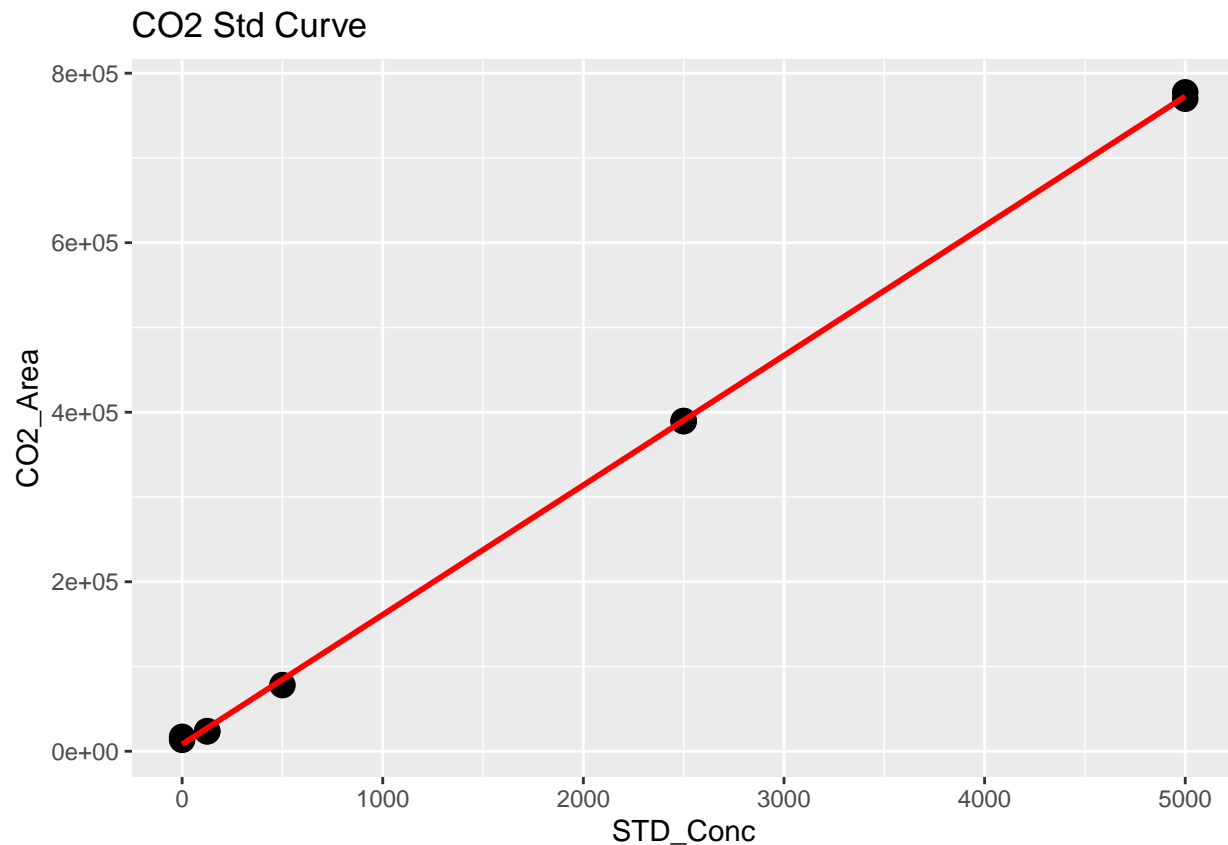
## 'geom_smooth()' using formula = 'y ~ x'
```

CH4 HIGH Std Curve



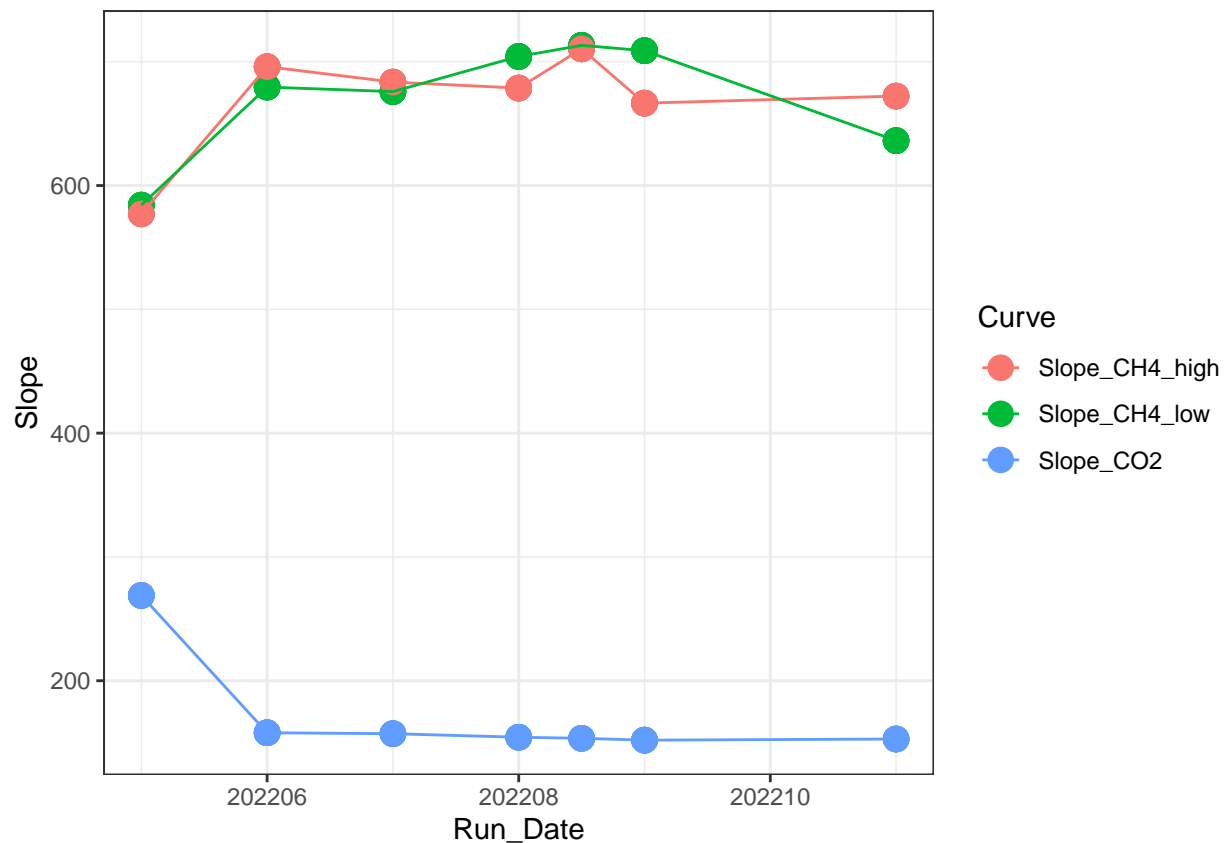
```
##
## Call:
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -28938  -6190  -4585       69   88709
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    5220.127    6726.336   0.776   0.449
## stds_ch4$STD_Conc  672.248      3.591 187.200 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 24710 on 16 degrees of freedom
## Multiple R-squared:  0.9995, Adjusted R-squared:  0.9995
## F-statistic: 3.504e+04 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
## -6593.6  -3598.0  -961.4   4658.3   8811.9
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    8210.1166   2428.5240     3.381   0.0117 *
## stds_co2$STD_Conc  152.9109     0.9195  166.299  7.5e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 5457 on 7 degrees of freedom
## Multiple R-squared:  0.9997, Adjusted R-squared:  0.9997
## F-statistic: 2.766e+04 on 1 and 7 DF, p-value: 7.502e-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)

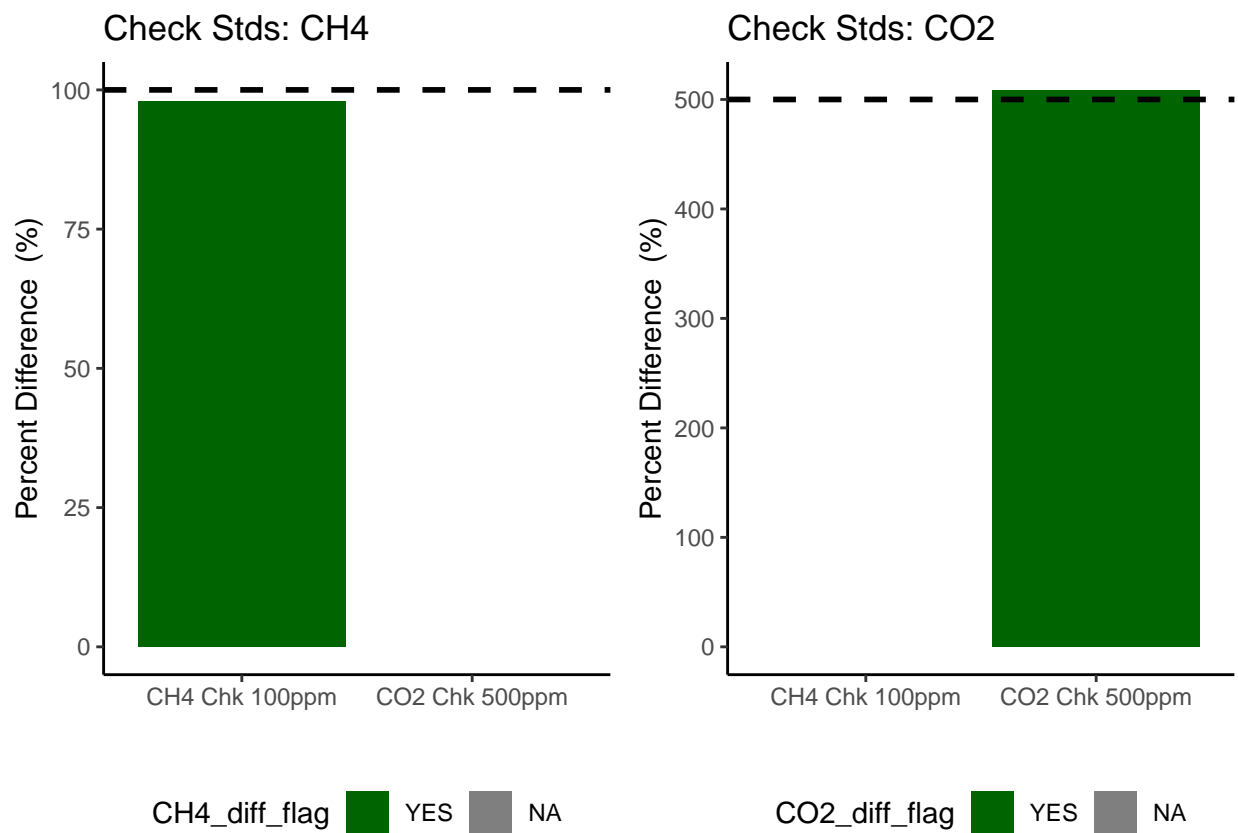
#####make flags for any dilutions needed
#highest CH4 standard = 10000
#highest CO2 standard = 50000

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.
```

```
## Warning: Removed 1 row containing missing values or values outside the scale range
## ('geom_bar()').
## Removed 1 row 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)

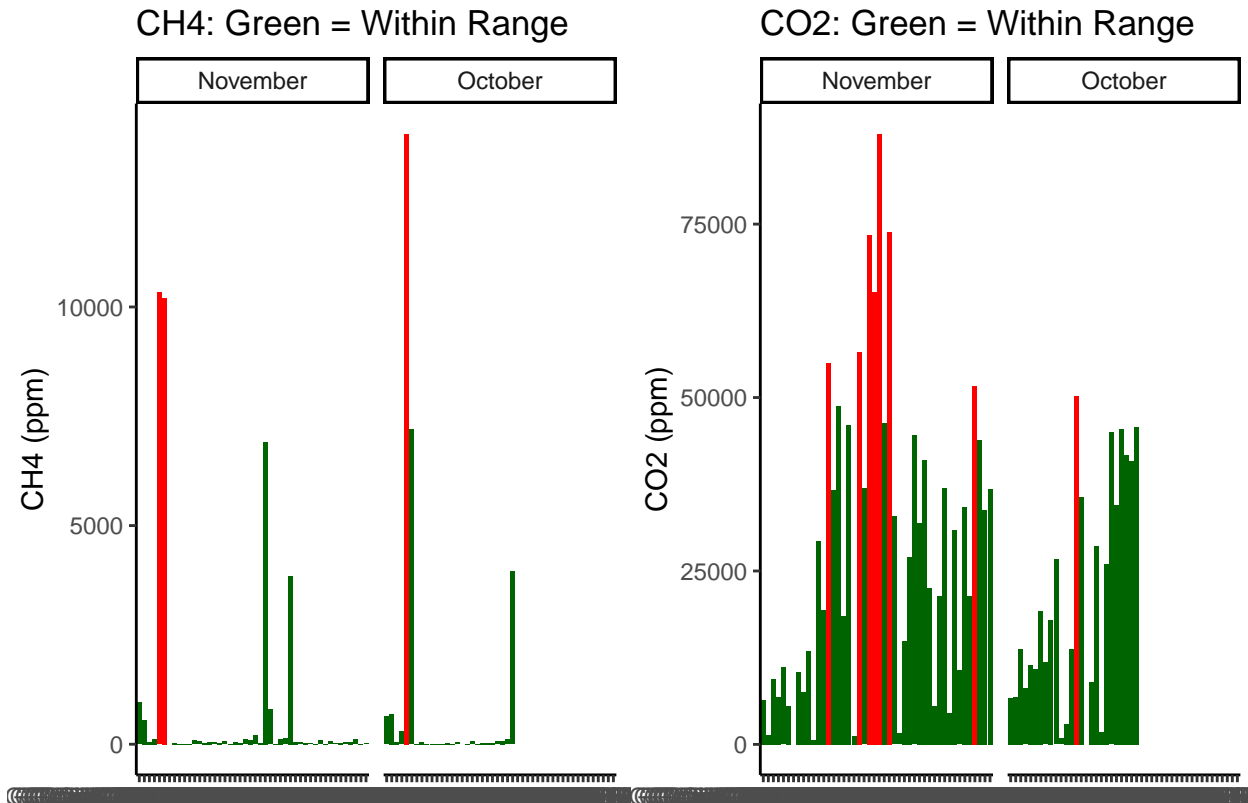
Samples$CO2_Conc_ppm_dilcorr <- (Samples$CO2_Conc_ppm * Samples$Dilution_Factor)

#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 202212.5   Unknown TGAS      2022
## 2 Varian GC Stephanie J. Wilson 202212.5   Unknown TGAS      2022
## 3 Varian GC Stephanie J. Wilson 202212.5   Unknown TGAS      2022
## 4 Varian GC Stephanie J. Wilson 202212.5   Unknown TGAS      2022
## 5 Varian GC Stephanie J. Wilson 202212.5   Unknown TGAS      2022
## 6 Varian GC Stephanie J. Wilson 202212.5   Unknown TGAS      2022
## Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      October GWI_TGAS_UP_SF_1              2      NA    18193 3989643
## 2      October GWI_TGAS_UP_SF_2              2      NA    22122 6882701
## 3      October GWI_TGAS_UP_SF_3              2      NA    21739 5279576
## 4      October GWI_TGAS_UP_SF_4              2      NA    45319 6962757
## 5      October GWI_TGAS_UP_SF_5              2      NA    52221 6386926
## 6      October GWI_TGAS_UP_SF_6              2      NA    80365 6263992
## Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm      CH4_Flag
## 1      NA      NA      Low    24.04512    26037.60 Within Range
## 2      NA      NA      Low    30.21995    44957.49 Within Range
## 3      NA      NA      Low    29.61803    34473.44 Within Range
## 4      NA      NA      Low    66.67646    45481.04 Within Range
```



```
## 5      NA      NA      Low      77.52367      41715.24 Within Range
## 6      NA      NA      High     111.78146      40911.28 Within Range
##      CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range      48.09024      52075.19
## 2 Within Range      60.43991      89914.98
## 3 Within Range      59.23606      68946.89
## 4 Within Range     133.35292      90962.07
## 5 Within Range     155.04734      83430.48
## 6 Within Range     223.56292      81822.57
```

```
#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 202212.5 TGAS      2022      October
## 2 Varian GC Stephanie J. Wilson 202212.5 TGAS      2022      October
## 3 Varian GC Stephanie J. Wilson 202212.5 TGAS      2022      October
## 4 Varian GC Stephanie J. Wilson 202212.5 TGAS      2022      October
## 5 Varian GC Stephanie J. Wilson 202212.5 TGAS      2022      October
## 6 Varian GC Stephanie J. Wilson 202212.5 TGAS      2022      October
##      Sample_ID Dilution_Factor Field_Notes      CH4_Flag      CO2_Flag
## 1 GWI_TGAS_UP_SF_1      2      NA Within Range Within Range
## 2 GWI_TGAS_UP_SF_2      2      NA Within Range Within Range
## 3 GWI_TGAS_UP_SF_3      2      NA Within Range Within Range
## 4 GWI_TGAS_UP_SF_4      2      NA Within Range Within Range
## 5 GWI_TGAS_UP_SF_5      2      NA Within Range Within Range
## 6 GWI_TGAS_UP_SF_6      2      NA Within Range Within Range
##      CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1      48.09024      52075.19
## 2      60.43991      89914.98
## 3      59.23606      68946.89
## 4     133.35292      90962.07
## 5     155.04734      83430.48
## 6     223.56292      81822.57
```

```
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(Samples1, "Processed Data/COMPASS_Synoptic_TGW_202210-11_Processed.csv")

```

#end