

# COMPASS\_Synoptic\_TGW\_2022: July

Stephanie J. Wilson

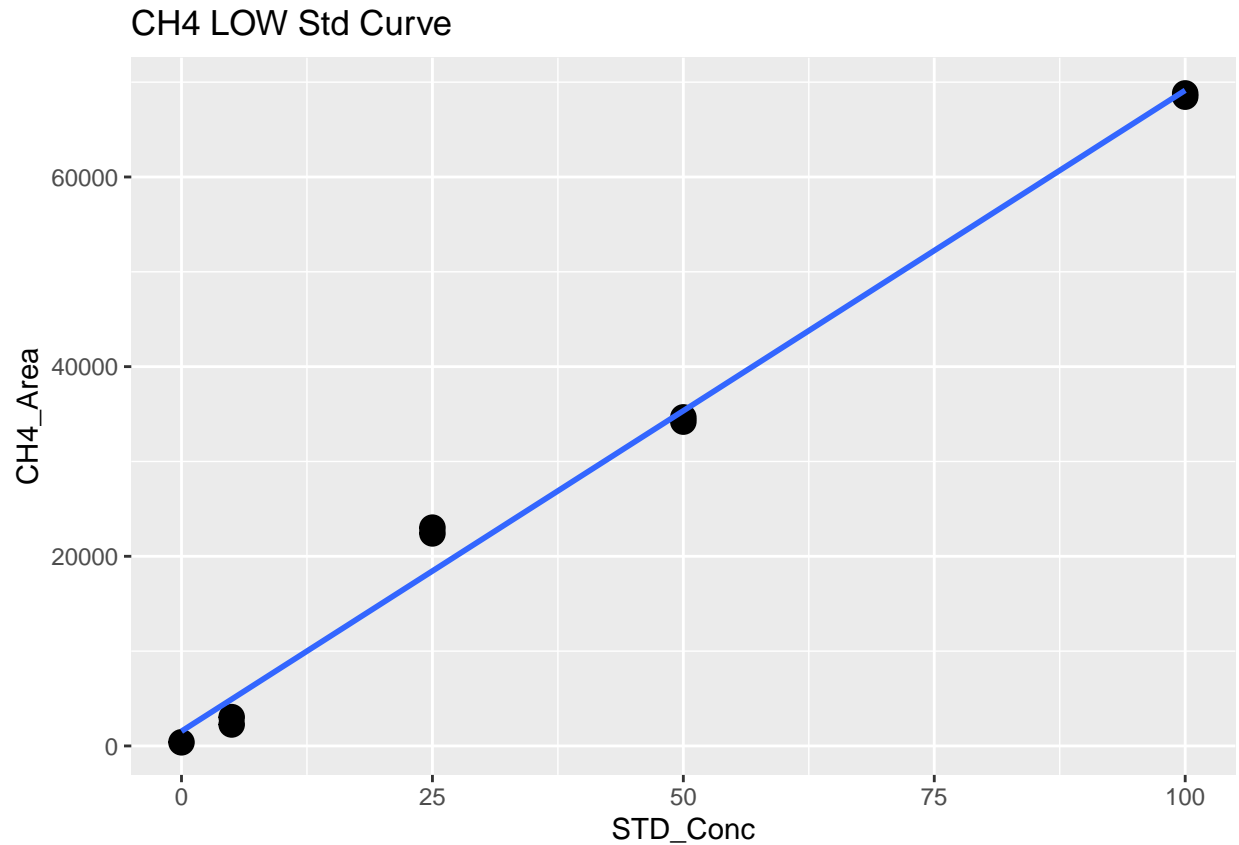
2023-01-14

##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 202207 Unknown TGAS 2022
## 2 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 3 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 4 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 5 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 6 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      July GCrew_Tgas_TR_SF_1      7      NA 387703 1228239
## 2      July GCrew_Tgas_TR_SF_2      7      NA 813894 1743563
## 3      July GCrew_Tgas_TR_SF_3      7      NA 98387 2163859
## 4      July GCrew_Tgas_TR_SF_4      7      NA 599507 2555958
## 5      July GCrew_Tgas_TR_SF_5      7      NA 8757124 1752010
## 6      July GCrew_Tgas_TR_SF_6      7      NA 13718650 1685685
##      Field.Notes Lab.Notes
## 1      NA      NA
## 2      NA      NA
## 3      NA      NA
## 4      NA      NA
## 5      NA      NA
## 6      NA      NA

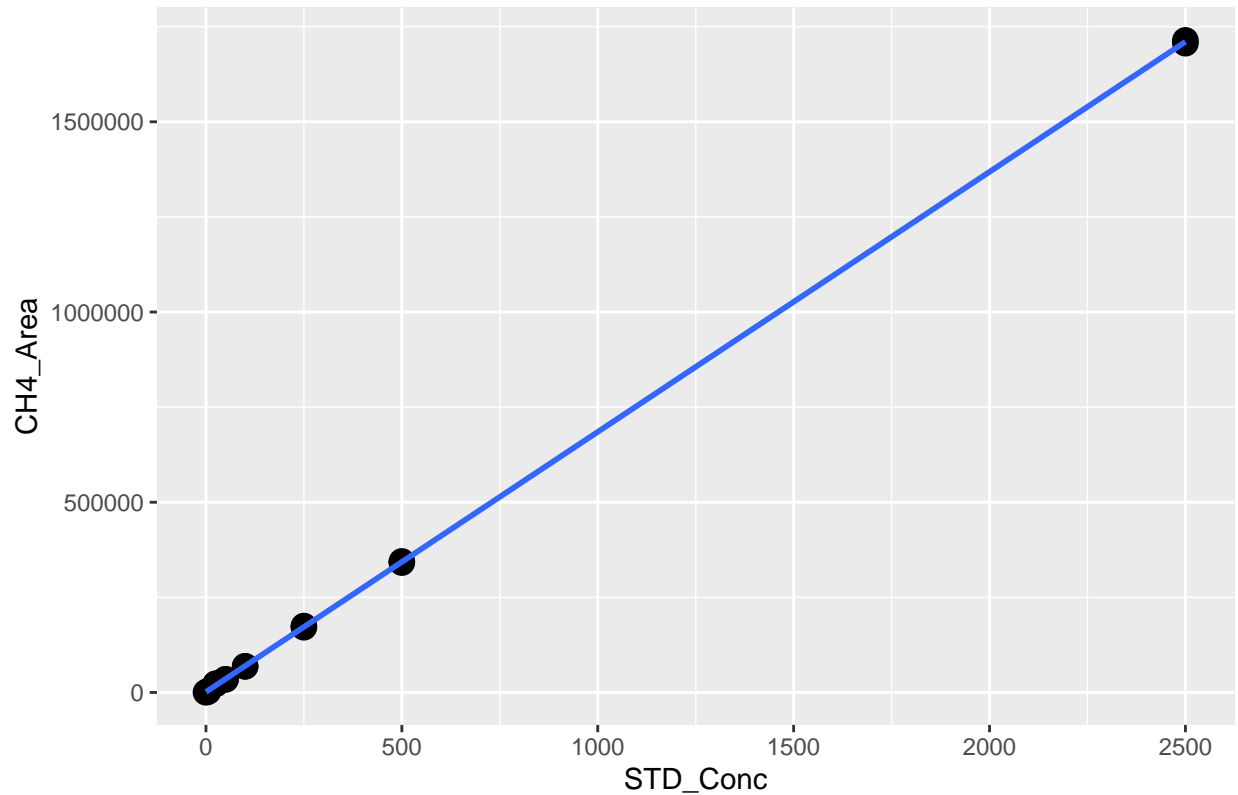
## 'geom_smooth()' using formula = 'y ~ x'
```



```
##
## Call:
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2663.1 -1163.6  -695.8  -324.1   4574.8
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1547.59    1338.16   1.157   0.285
## stds_ch4_low$STD_Conc    675.90     24.75  27.304 2.27e-08 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2700 on 7 degrees of freedom
## Multiple R-squared:  0.9907, Adjusted R-squared:  0.9894
## F-statistic: 745.5 on 1 and 7 DF, p-value: 2.267e-08

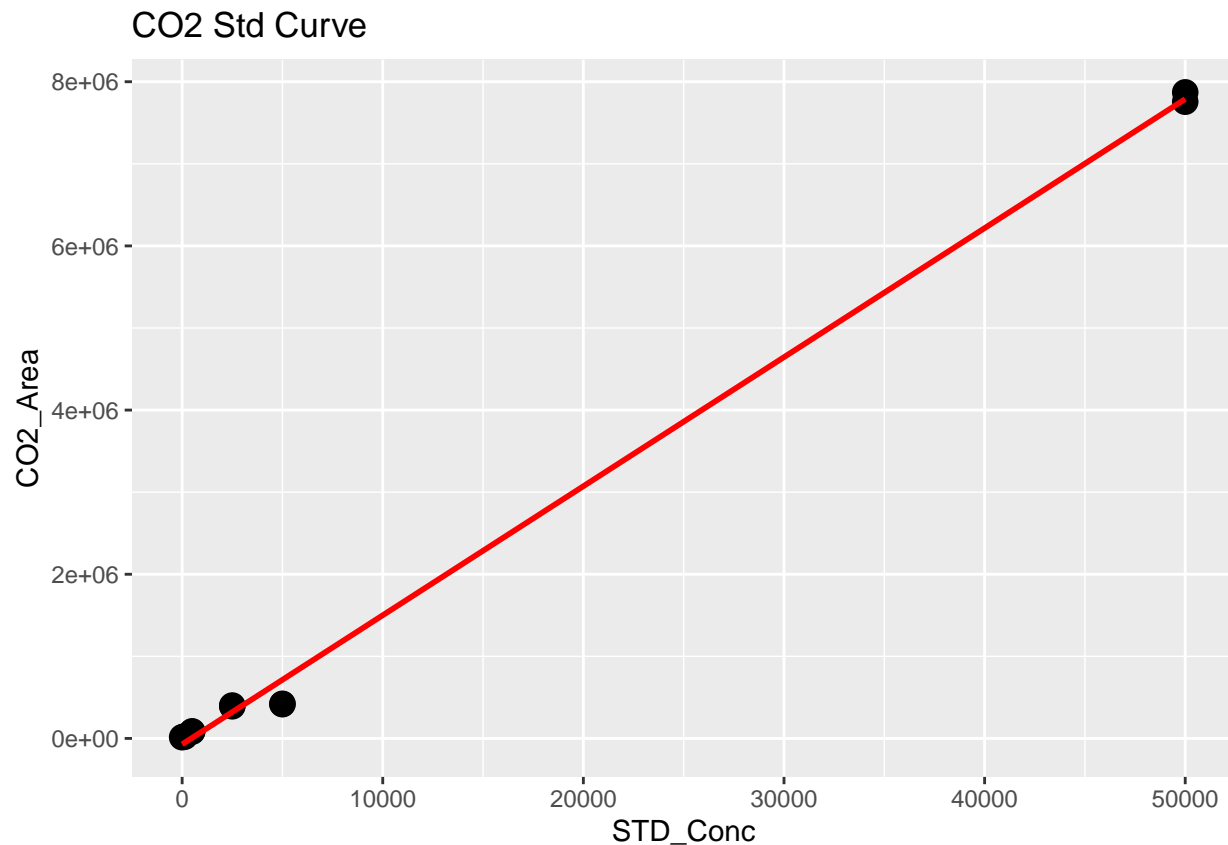
## 'geom_smooth()' using formula = 'y ~ x'
```

CH4 HIGH Std Curve



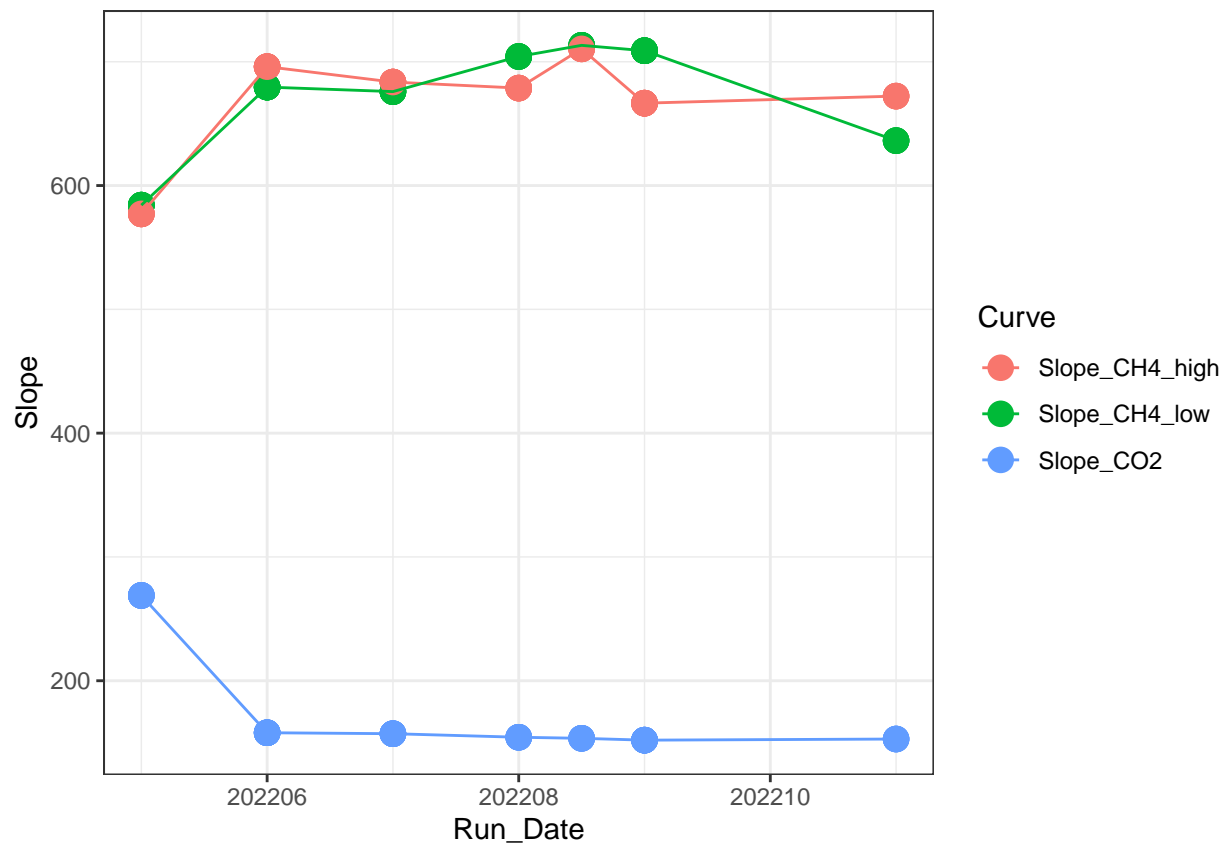
```
##
## Call:
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -3795.6 -1512.2  -948.6   1849.4   4597.7
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1332.5553    790.6757   1.685   0.116
## stds_ch4$STD_Conc  683.5880     0.8444 809.533 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2672 on 13 degrees of freedom
## Multiple R-squared:  1, Adjusted R-squared:  1
## F-statistic: 6.553e+05 on 1 and 13 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
## -297324   18739   75227   78479   86329
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -71502.431   54643.174  -1.309    0.223
## stds_co2$STD_Conc    157.207     2.547   61.723 3.88e-13 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 157700 on 9 degrees of freedom
## Multiple R-squared:  0.9976, Adjusted R-squared:  0.9974
## F-statistic: 3810 on 1 and 9 DF, p-value: 3.879e-13

##   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
```

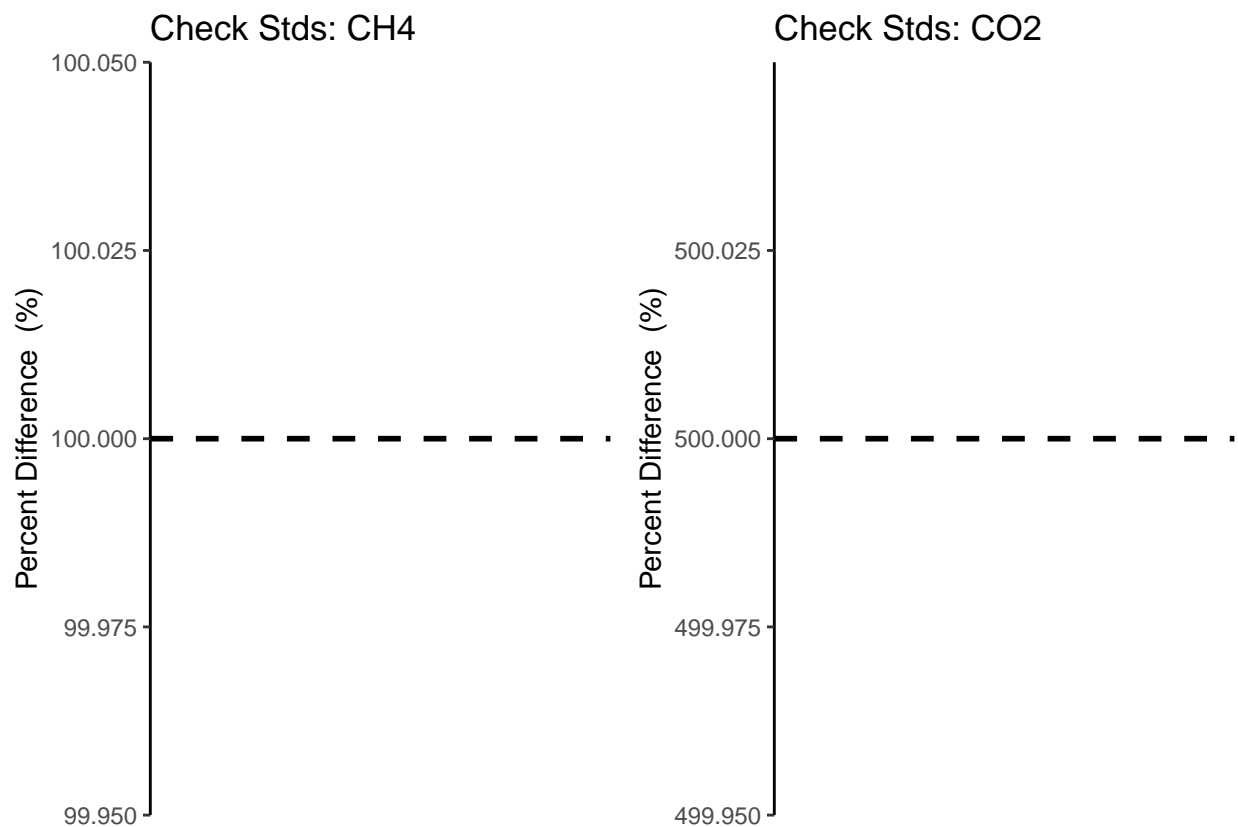
```
#head(Samples)

#####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)

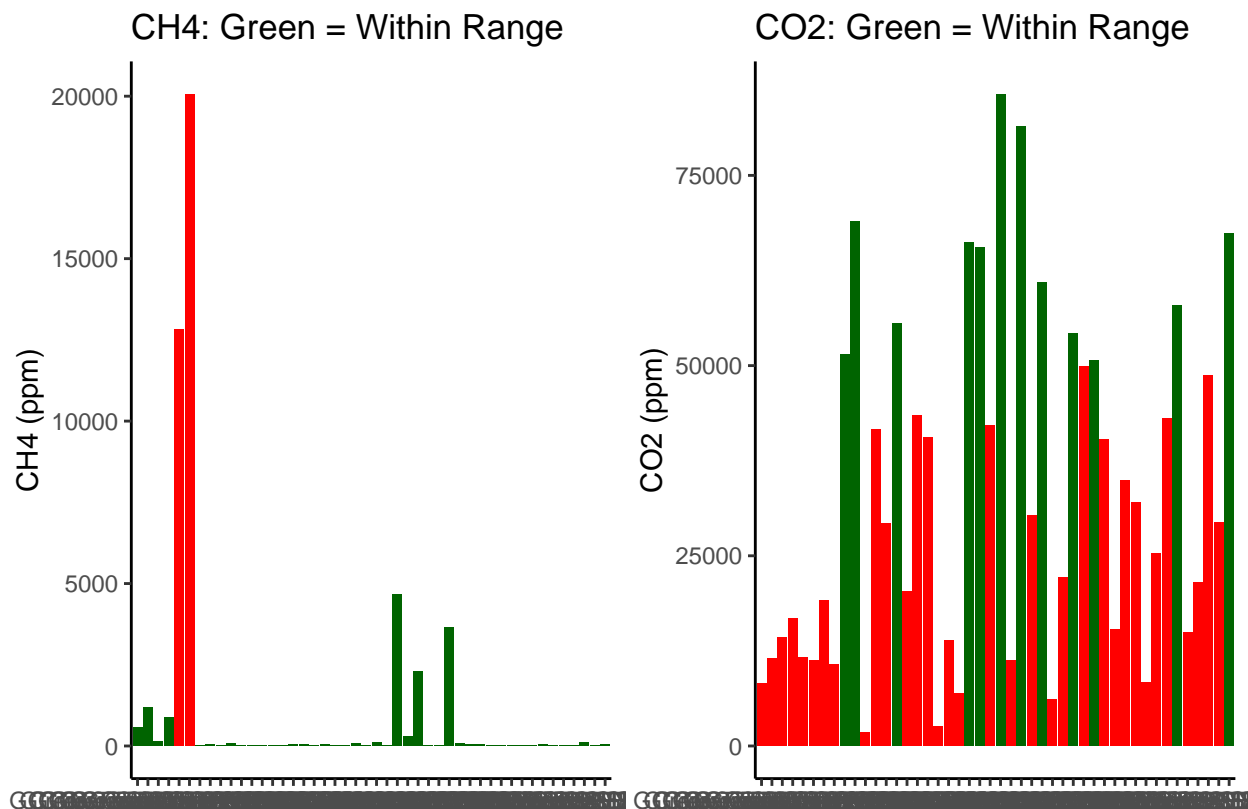
#check results
#head(Samples)

#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( "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")

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")

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 202207 Unknown TGAS 2022
## 2 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 3 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 4 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 5 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
## 6 Varian GC Stephanie J. Wilson 202207 Unknown TGAS 2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1 July GCrew_Tgas_TR_SF_1 7 NA 387703 1228239
## 2 July GCrew_Tgas_TR_SF_2 7 NA 813894 1743563
## 3 July GCrew_Tgas_TR_SF_3 7 NA 98387 2163859
## 4 July GCrew_Tgas_TR_SF_4 7 NA 599507 2555958
## 5 July GCrew_Tgas_TR_SF_5 7 NA 8757124 1752010
## 6 July GCrew_Tgas_TR_SF_6 7 NA 13718650 1685685
##      Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm CH4_Flag
## 1 NA NA High 565.2095 8267.732 Within Range
## 2 NA NA High 1188.6713 11545.738 Within Range
## 3 NA NA High 141.9780 14219.266 Within Range
## 4 NA NA High 875.0511 16713.431 Within Range
## 5 NA NA High 12808.5798 11599.470 Needs Dilution
## 6 NA NA High 20066.6446 11177.573 Needs Dilution
##      CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range 3956.4666 57874.12
## 2 Within Range 8320.6992 80820.17
## 3 Within Range 993.8459 99534.86
## 4 Within Range 6125.3578 116994.02
## 5 Within Range 89660.0588 81196.29
## 6 Within Range 140466.5125 78243.01
```

```
#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 202207 TGAS 2022 July
## 2 Varian GC Stephanie J. Wilson 202207 TGAS 2022 July
## 3 Varian GC Stephanie J. Wilson 202207 TGAS 2022 July
## 4 Varian GC Stephanie J. Wilson 202207 TGAS 2022 July
## 5 Varian GC Stephanie J. Wilson 202207 TGAS 2022 July
## 6 Varian GC Stephanie J. Wilson 202207 TGAS 2022 July
##      Sample_ID Dilution_Factor Field.Notes CH4_Flag CO2_Flag
## 1 GCrew_Tgas_TR_SF_1 7 NA Within Range Within Range
## 2 GCrew_Tgas_TR_SF_2 7 NA Within Range Within Range
## 3 GCrew_Tgas_TR_SF_3 7 NA Within Range Within Range
## 4 GCrew_Tgas_TR_SF_4 7 NA Within Range Within Range
## 5 GCrew_Tgas_TR_SF_5 7 NA Needs Dilution Within Range
## 6 GCrew_Tgas_TR_SF_6 7 NA Needs Dilution Within Range
```



```
##   CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1           3956.4666           57874.12
## 2           8320.6992           80820.17
## 3           993.8459           99534.86
## 4           6125.3578          116994.02
## 5           89660.0588          81196.29
## 6          140466.5125          78243.01
```

```
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_202207_Processed.csv")
```

```
#end
```