

COMPASS_Synoptic_SGW_2023: November

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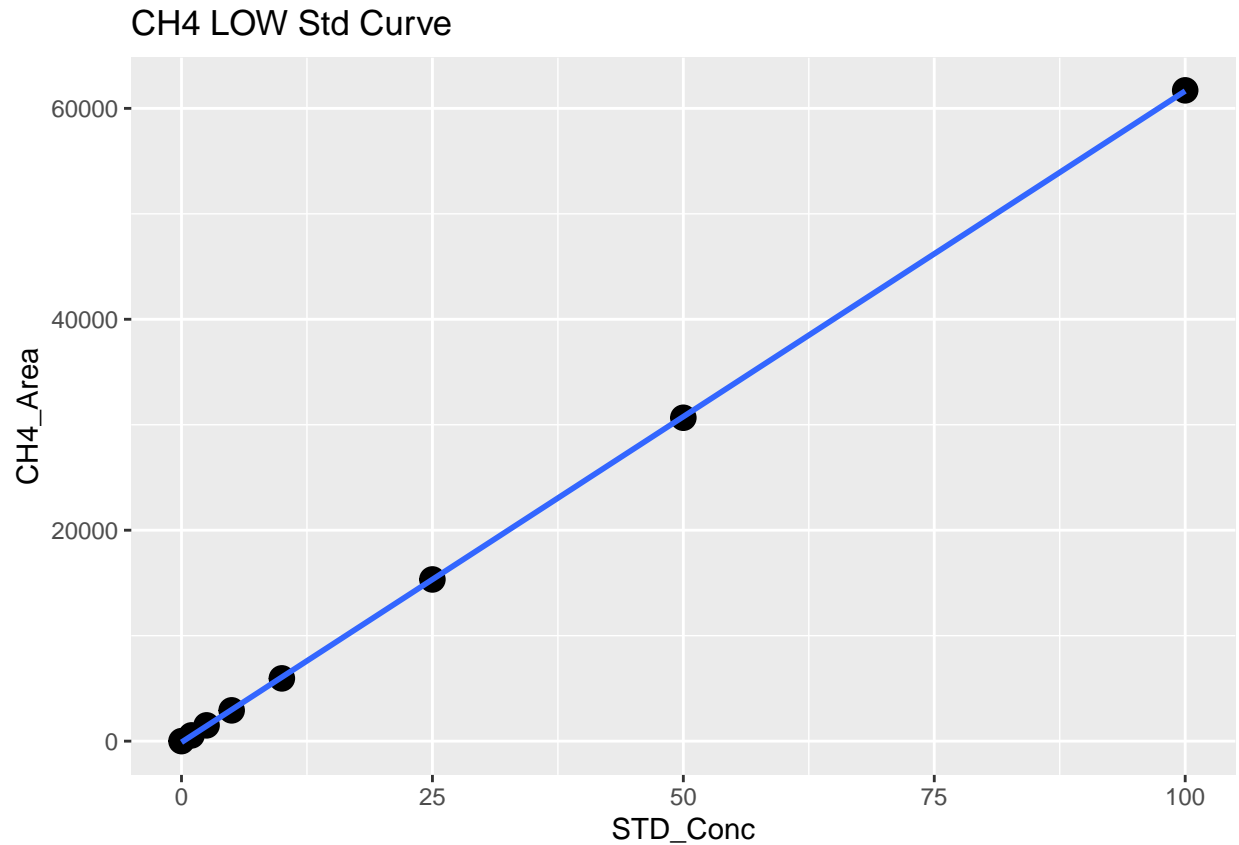
##Set Up

Read in first data file and assess standard curves

##	Machine	User	Run_Date	Sample_Type	Type1	Sample_Year	Sample_Month
## 1	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>
## 2	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>
## 3	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>
## 4	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>
## 5	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>
## 6	Varian GC	Wegner	11/28/2023	STD_CH4	STDs	2023	<NA>

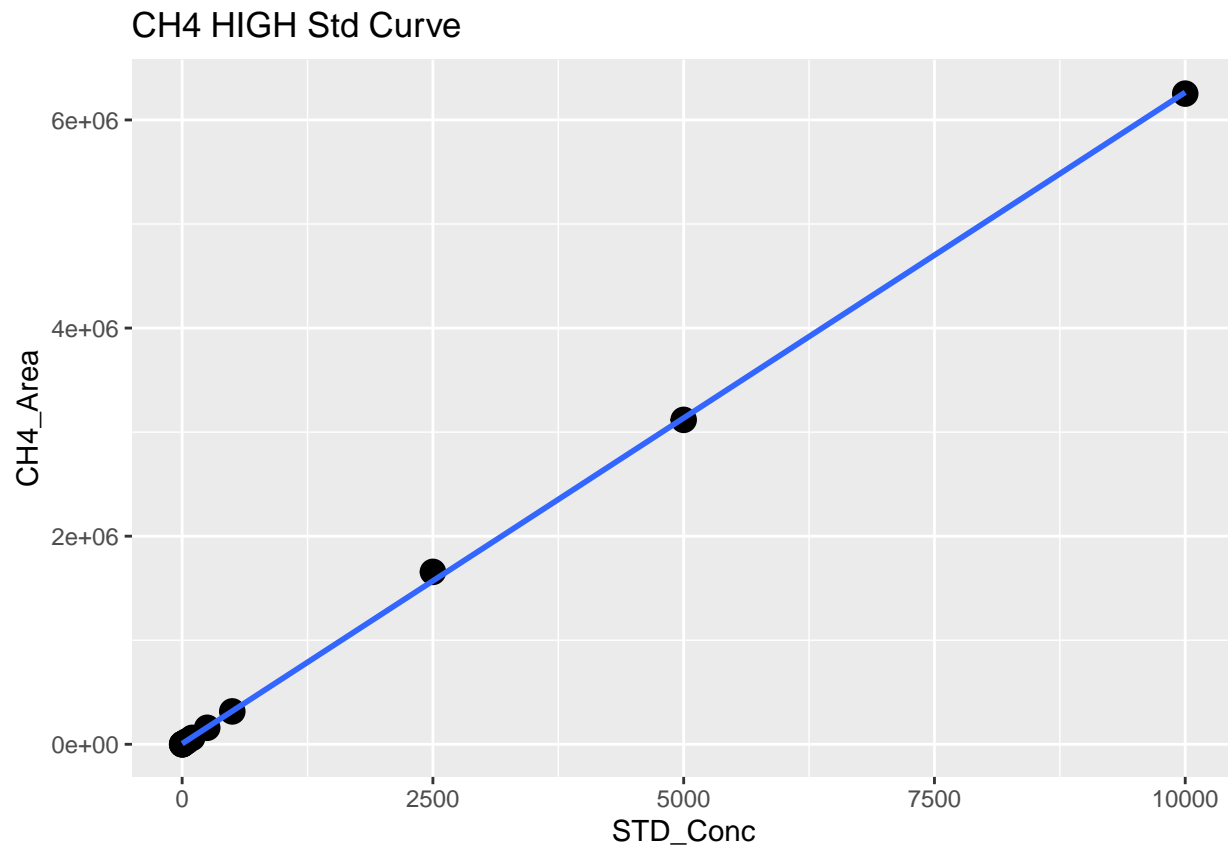
##	Sample_ID	Dilution_Factor	STD_Conc	CO2_Area	CH4_Area	Lab_Notes
## 1	CH4	1	0.0	0	0	NA
## 2	CH4	1	1.0	19812	545	NA
## 3	CH4	1	2.5	65112	1497	NA
## 4	CH4	1	5.0	130478	2919	NA
## 5	CH4	1	10.0	266905	5951	NA
## 6	CH4	1	25.0	0	15332	NA

'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
## -114.49  -70.19   20.36   63.33  110.02
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    -110.0215     39.7533  -2.768   0.0325 *
## stds_ch4_low$STD_Conc  617.5513     0.9765 632.385 1.06e-15 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 90.44 on 6 degrees of freedom
## Multiple R-squared:  1, Adjusted R-squared:  1
## F-statistic: 3.999e+05 on 1 and 6 DF, p-value: 1.055e-15

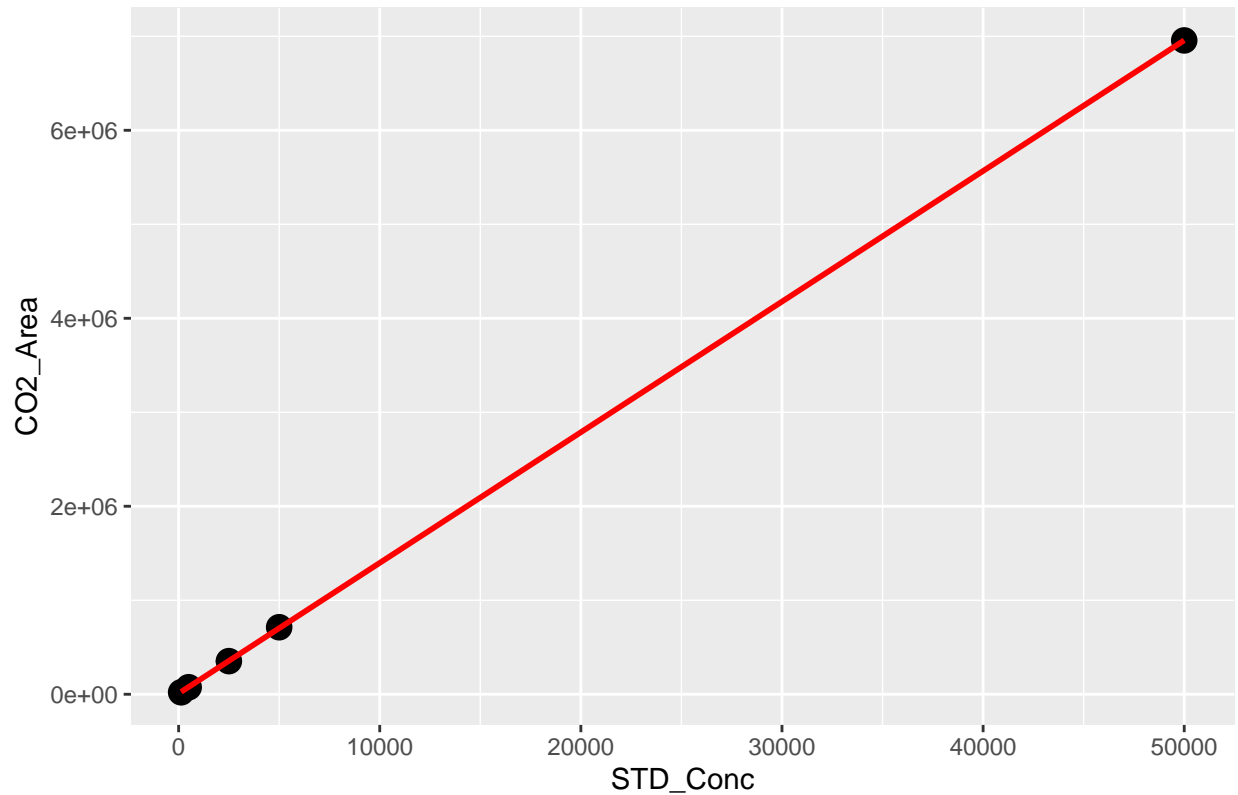
## 'geom_smooth()' using formula = 'y ~ x'
```



```
##
## Call:
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -17758  -6526  -6102  -5891   86227
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    5891.337    8477.033   0.695   0.501
## stds_ch4$STD_Conc  625.944      2.665 234.913 <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 27360 on 11 degrees of freedom
## Multiple R-squared:  0.9998, Adjusted R-squared:  0.9998
## F-statistic: 5.518e+04 on 1 and 11 DF, p-value: < 2.2e-16

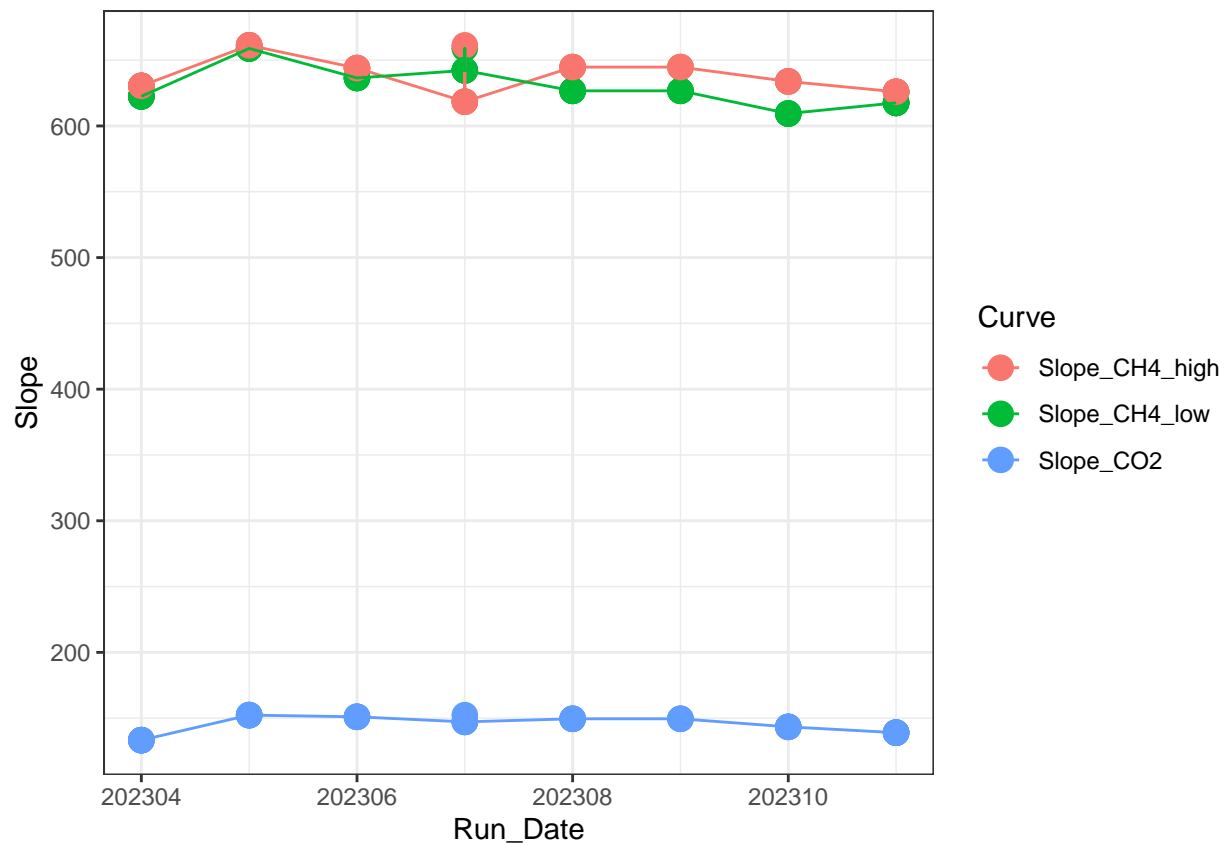
## 'geom_smooth()' using formula = 'y ~ x'
```

CO2 Std Curve



```
##
## Call:
## lm(formula = stds_co2$CO2_Area ~ stds_co2$STD_Conc)
##
## Residuals:
##      1      2      3      4      5
## -4255 -2474 -2785 10376  -863
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    7748.1692   3572.9453     2.169   0.119
## stds_co2$STD_Conc 138.9870     0.1588  875.292 3.29e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 6841 on 3 degrees of freedom
## Multiple R-squared:      1, Adjusted R-squared:      1
## F-statistic: 7.661e+05 on 1 and 3 DF, p-value: 3.289e-09

##      X      Curve      R2      Slope      Intercept Run_Date
## 1 1  Slope_CH4_low 0.9997982 642.2013   -176.61917  202307
## 2 2  Slope_CH4_high 0.9995317 618.4551  11202.40792  202307
## 3 3      Slope_CO2 0.9999842 147.1737  10330.39781  202307
## 4 4  Slope_CH4_low 0.9990979 658.8841   -87.56451  202307
## 5 5  Slope_CH4_high 0.9999901 661.2523   176.62351  202307
## 6 6      Slope_CO2 0.9999978 152.3154   8787.66721  202307
```



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, "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)
```

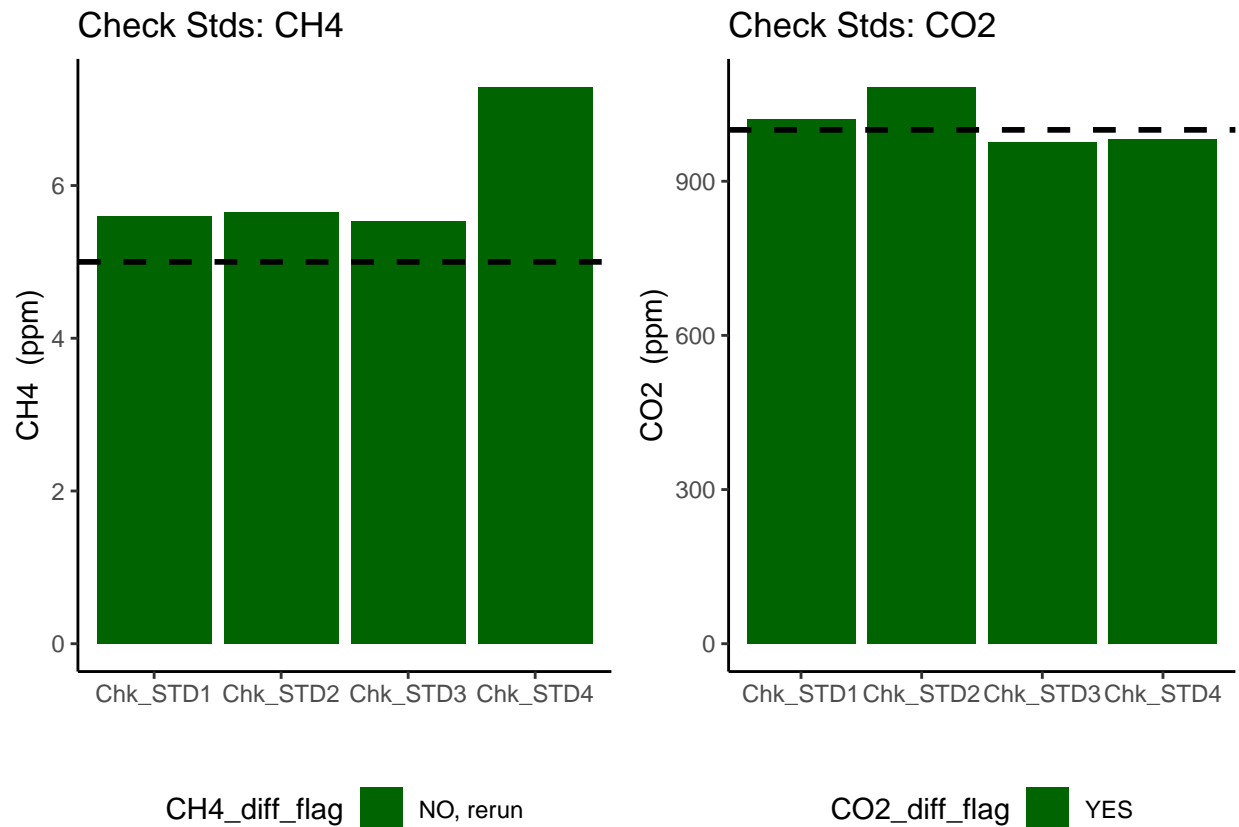
```
#head(Samples)

#####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)

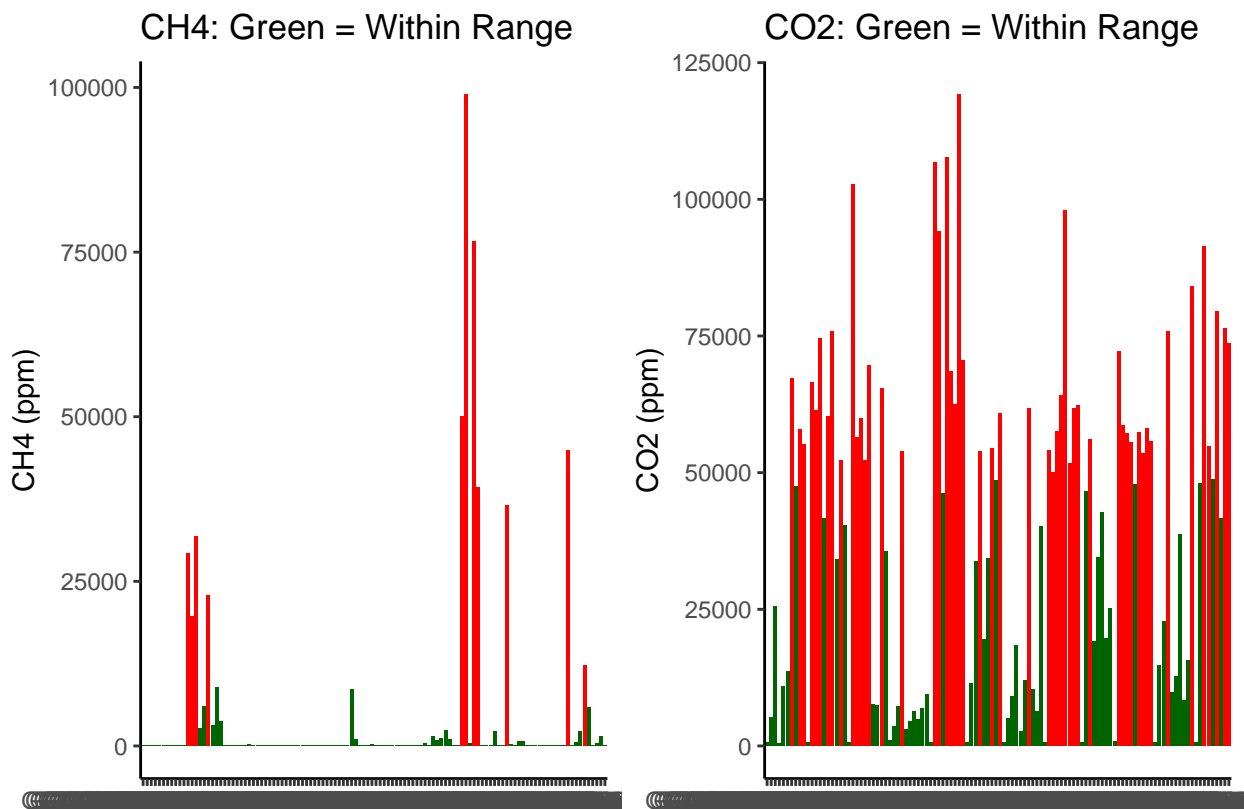
#check results
#head(Samples)

#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("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)

```



If samples are water calculate gas in water - only need if there is water

Write out processed data & slopes

```
#pull out what we need
Samples1 <- Samples[,c(1:3,6:9,17:20)]
#head(Samples1)

IDs <- data.frame(do.call('rbind', strsplit(as.character(Samples1$Sample_ID), '_ ', fixed=TRUE)))

## Warning in rbind(c("MSM", "UP", "sgwA", "10cm"), c("MSM", "UP", "SgwA", :
## number of columns of result is not a multiple of vector length (arg 10)

colnames(IDs) <- c("Site", "Zone", "Replicate", "Depth")
IDs$Depth1 <- ifelse(IDs$Depth == '10cm', '10',
                     ifelse(IDs$Depth == '20cm', '20',
                             ifelse(IDs$Depth == '45cm', '45', '0'))))
head(IDs)

##   Site Zone Replicate Depth Depth1
## 1  MSM   UP      sgwA  10cm     10
## 2  MSM   UP      SgwA  20cm     20
## 3  MSM   UP      SgwA  45cm     45
## 4  MSM   UP      SgwB  10cm     10
## 5  MSM   UP      SgwB  20cm     20
## 6  MSM   UP      SgwB  45cm     45

#rejoin them to the dataframe
alldat <- cbind(IDs, Samples1)
head(alldat)

##   Site Zone Replicate Depth Depth1 Machine User Run_Date Sample_Year
## 1  MSM   UP      sgwA  10cm     10 Varian GC Wegner 11/28/2023      2023
## 2  MSM   UP      SgwA  20cm     20 Varian GC Wegner 11/28/2023      2023
## 3  MSM   UP      SgwA  45cm     45 Varian GC Wegner 11/28/2023      2023
## 4  MSM   UP      SgwB  10cm     10 Varian GC Wegner 11/28/2023      2023
## 5  MSM   UP      SgwB  20cm     20 Varian GC Wegner 11/28/2023      2023
## 6  MSM   UP      SgwB  45cm     45 Varian GC Wegner 11/28/2023      2023
##   Sample_Month Sample_ID Dilution_Factor CH4_Flag CO2_Flag
## 1   November MSM_UP_sgwA_10cm           1 Within Range Within Range
## 2   November MSM_UP_SgwA_20cm           1 Within Range Within Range
## 3   November MSM_UP_SgwA_45cm           1 Within Range Within Range
## 4   November MSM_UP_SgwB_10cm           1 Within Range Within Range
## 5   November MSM_UP_SgwB_20cm           1 Within Range Within Range
## 6   November MSM_UP_SgwB_45cm           1 Within Range Needs Dilution
##   CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1             3.611071             4963.773
## 2             3.376273             9050.759
## 3             3.054032            18396.468
## 4             3.141474             2628.489
## 5             2.961732            11943.246
## 6             2.454892            61812.695
```



```
write.csv(alldat, "Processed Data/COMPASS_CBSYN_SGW_202311_Processed.csv")
```

```
#end
```