

COMPASS_Synoptic_TGW_2022: Sept & Oct part 1

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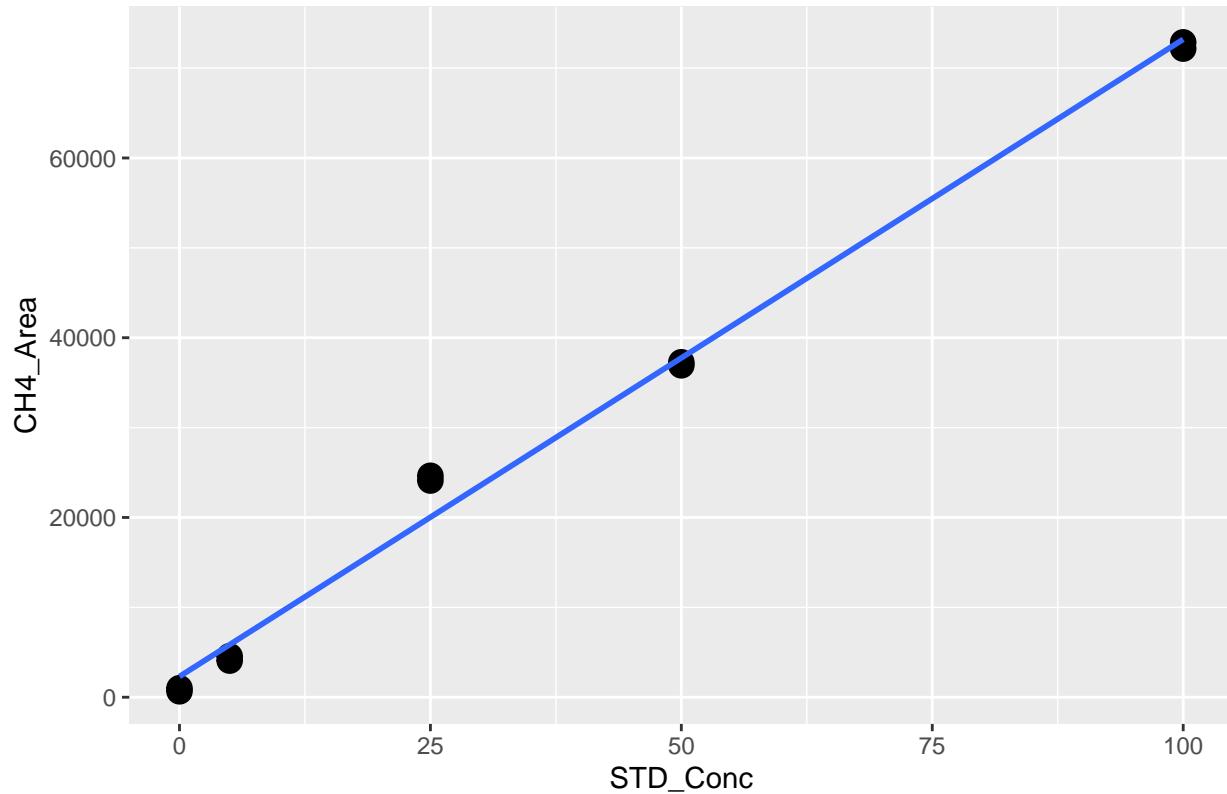
2023-01-14

##Set Up

Read in first data file and assess standard curves

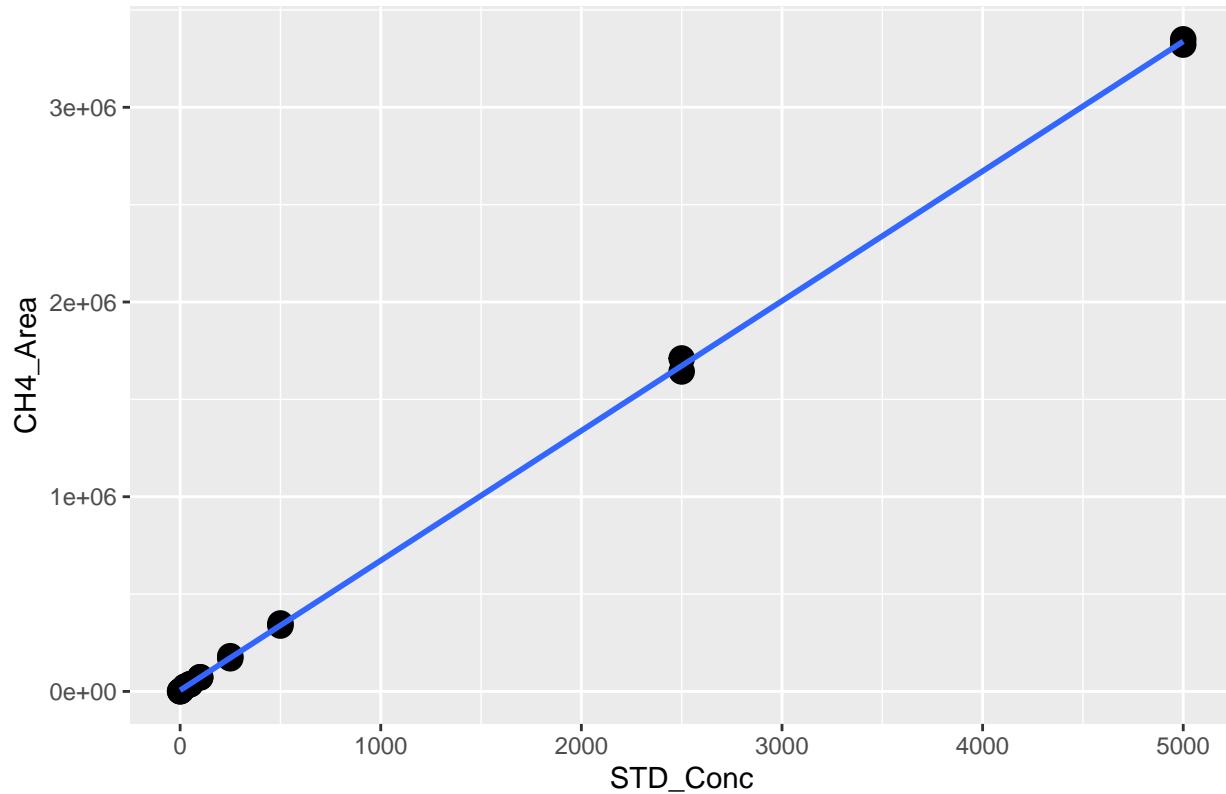
```
##      Machine          User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202212     Chk_STD  TGAS      2022
## 2 Varian GC Stephanie J. Wilson 202212     Chk_STD  TGAS      2022
## 3 Varian GC Stephanie J. Wilson 202212 Unknown  TGAS      2022
## 4 Varian GC Stephanie 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'
```

CH4 LOW Std Curve

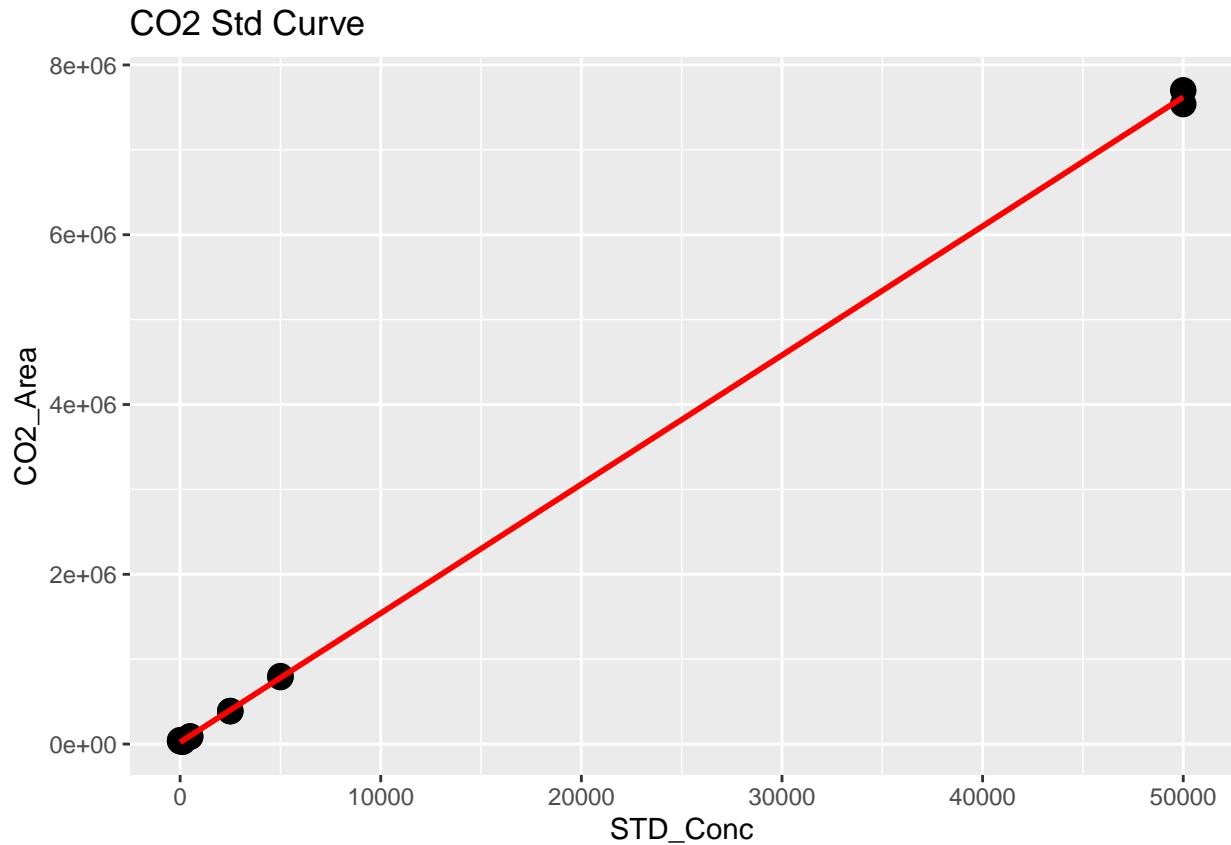


```
##  
## Call:  
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -17777.5 -1300.3  -928.8  -370.2  4611.0  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (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'
```

CH4 HIGH Std Curve

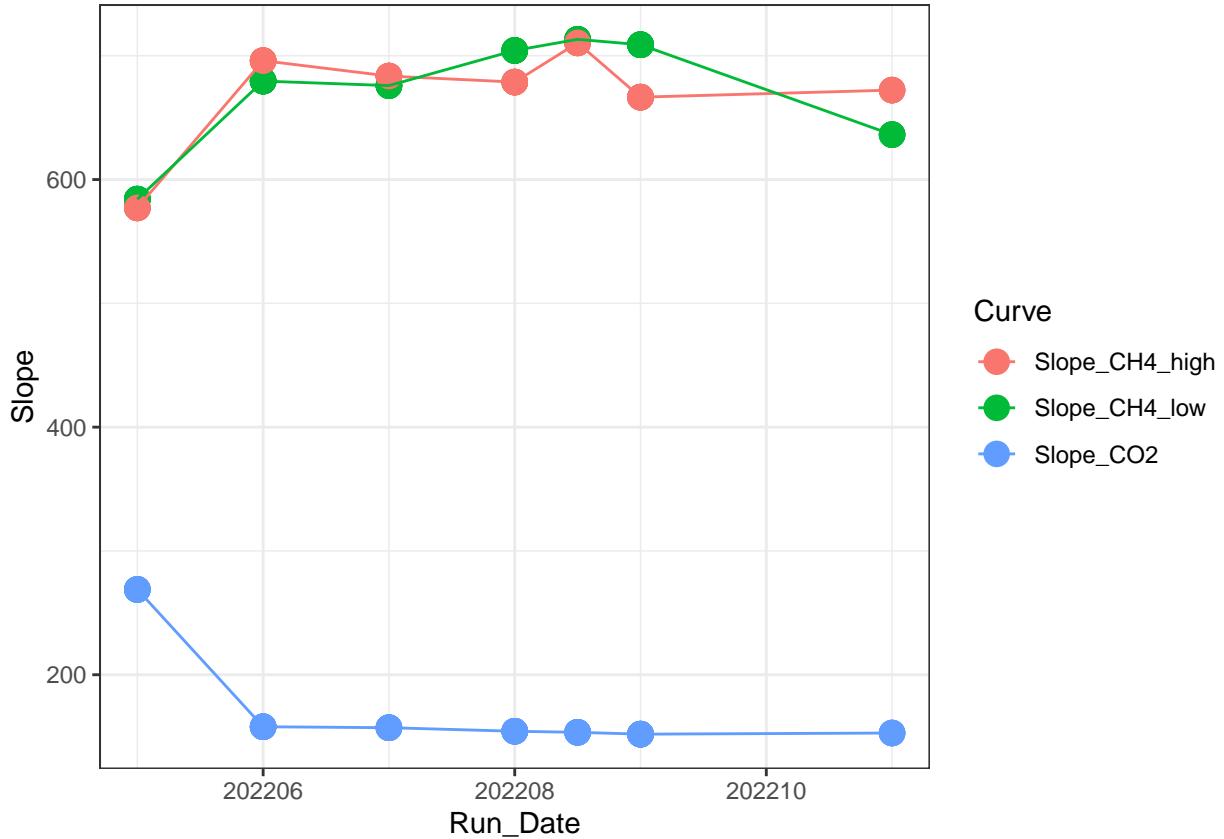


```
##  
## Call:  
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -28677  -4679  -1195   2106  37909  
##  
## Coefficients:  
##                 Estimate Std. Error t value Pr(>|t|)  
## (Intercept) 5738.283   3679.618   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)

```

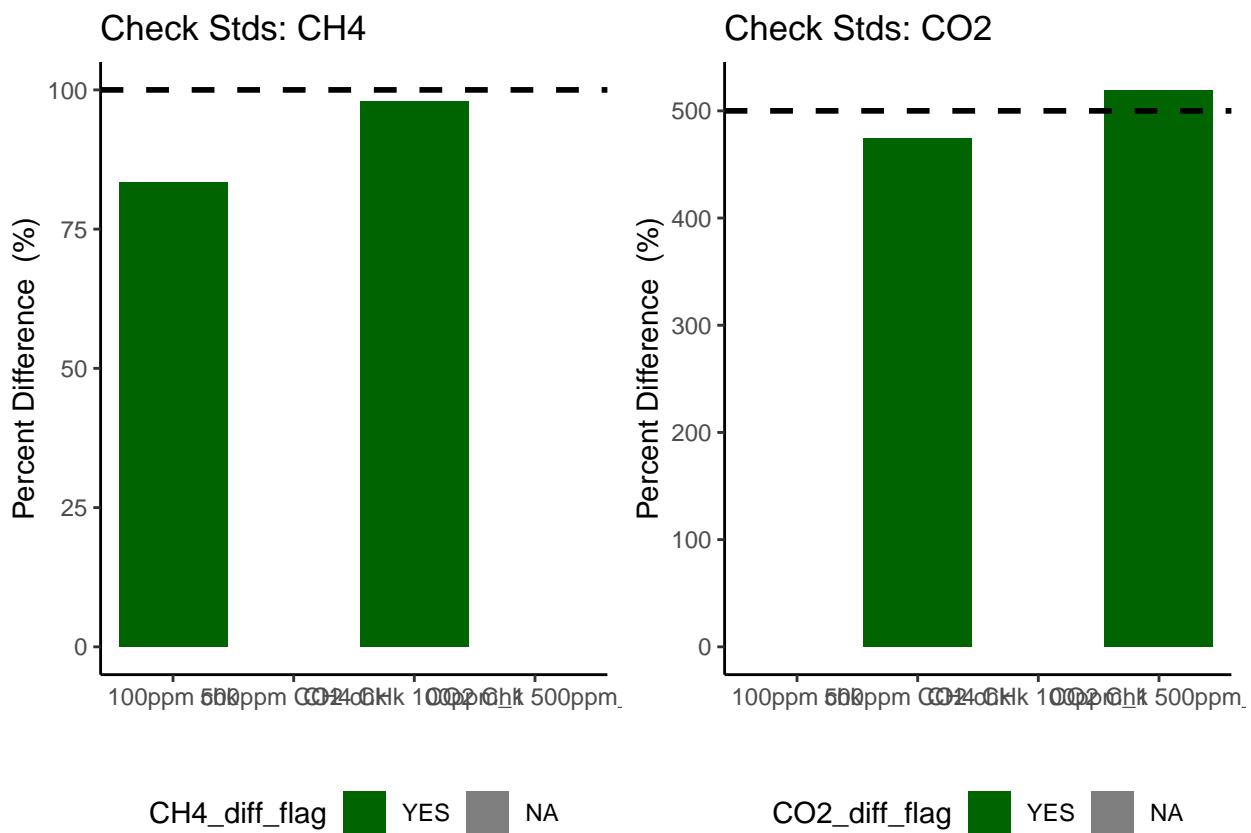
Check the Check Standards

```

## 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)

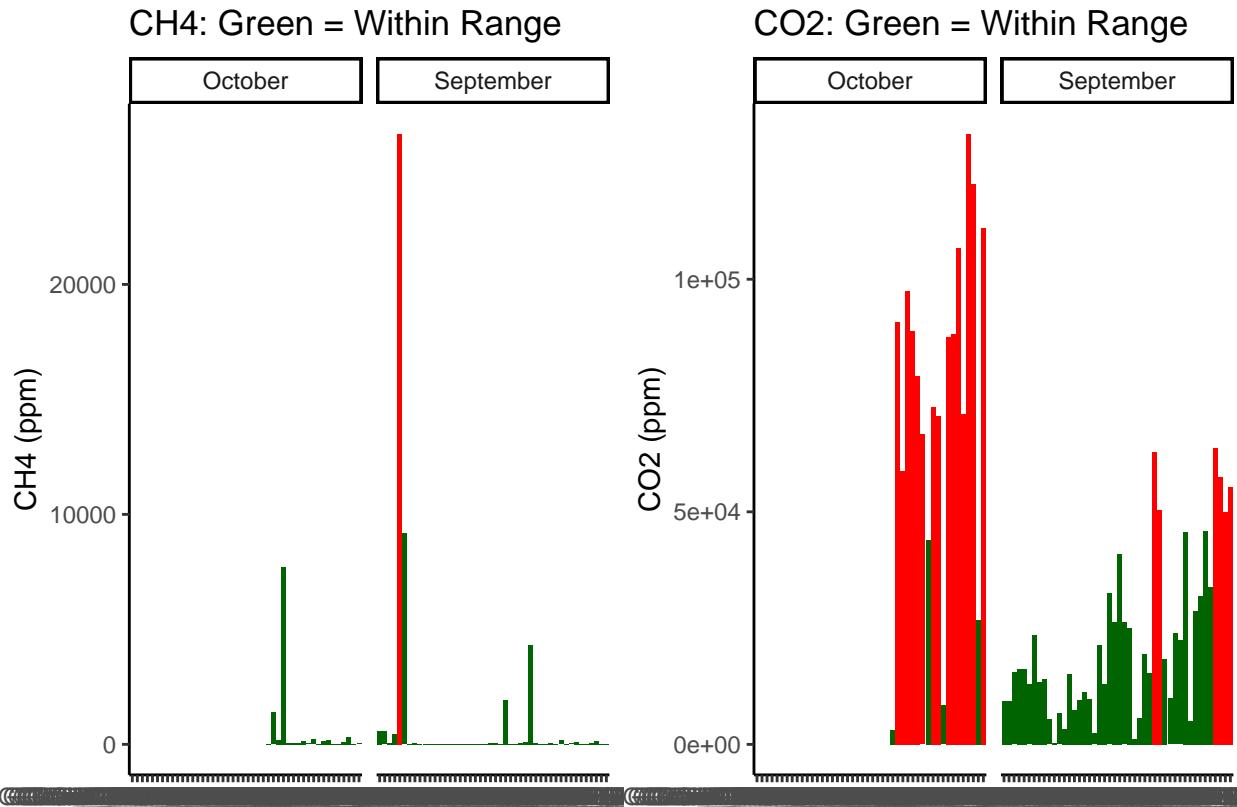
Samples$CO2_Conc_ppm_dilcorr <- (Samples$CO2_Conc_ppm * Samples$Dilution_Factor)

#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 = 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 ,
         Project = Project,
         Region = Region)

```

```

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