

COMPASS_Synoptic_TGW_2022: May

Stephanie J. Wilson

2023-01-14

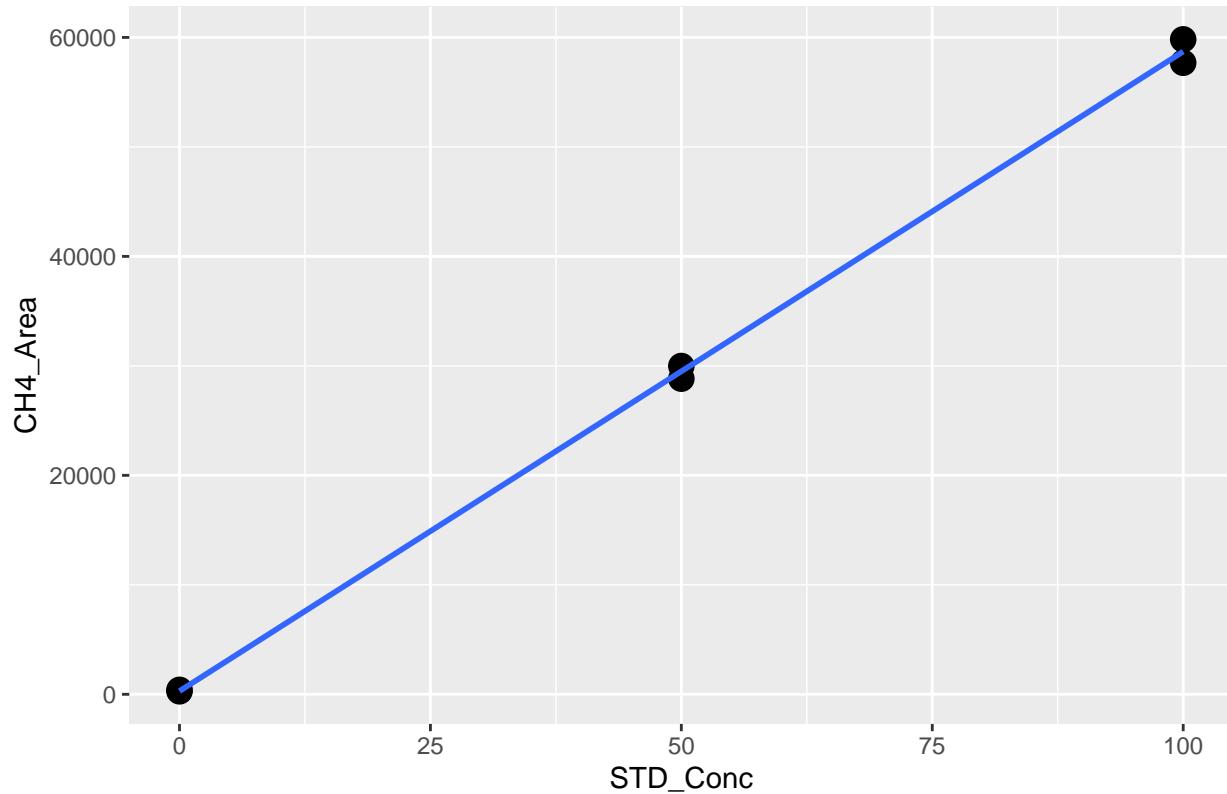
##Set Up

Read in first data file and assess standard curves

```
##      Machine          User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
## 2 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
## 3 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
## 4 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
## 5 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
## 6 Varian GC Stephanie J. Wilson 202205    Unknown  TGAS     2022
##   Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      May GWI_TGAS_TR_DS_1           2       NA    1420  285551
## 2      May GWI_TGAS_TR_DS_2           2       NA    5068 1114202
## 3      May GWI_TGAS_TR_DS_3           2       NA    2119 4462563
## 4      May GWI_TGAS_TR_SF_1           2       NA    1847 3899395
## 5      May GWI_TGAS_TR_SF_2           2       NA    1424  856806
## 6      May GWI_TGAS_TR_SF_3           2       NA    3065 2703534
##   Field.Notes Lab.Notes
## 1       NA       NA
## 2       NA       NA
## 3       NA       NA
## 4       NA       NA
## 5       NA       NA
## 6       NA       NA

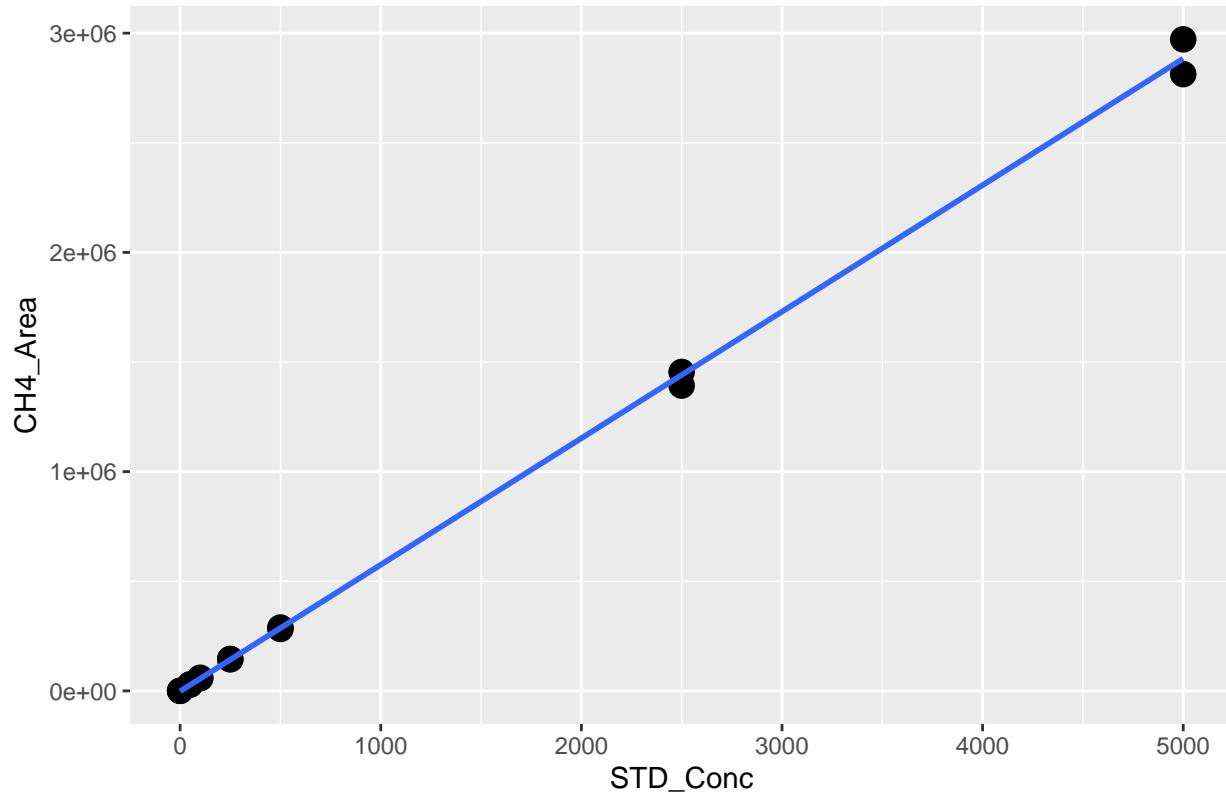
## `geom_smooth()` using formula = 'y ~ x'
```

CH4 LOW Std Curve



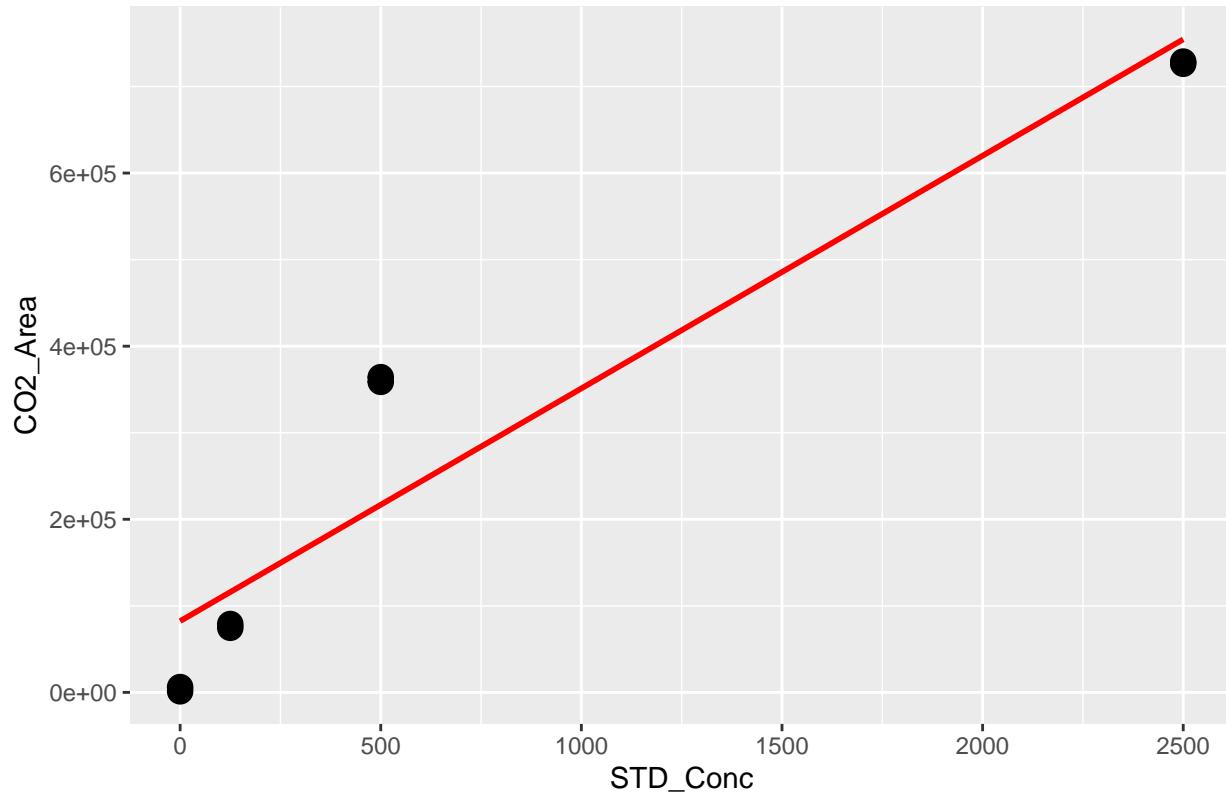
```
##  
## Call:  
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)  
##  
## Residuals:  
##      1       2       3       4       5       6  
## -18.67  112.33  484.83 -672.17 1116.33 -1022.67  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (Intercept)            292.667    558.260   0.524   0.628  
## stds_ch4_low$STD_Conc  584.170     8.649  67.546 2.88e-07 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 864.9 on 4 degrees of freedom  
## Multiple R-squared:  0.9991, Adjusted R-squared:  0.9989  
## F-statistic:  4562 on 1 and 4 DF,  p-value: 2.878e-07  
  
## 'geom_smooth()' using formula = 'y ~ x'
```

CH4 HIGH Std Curve



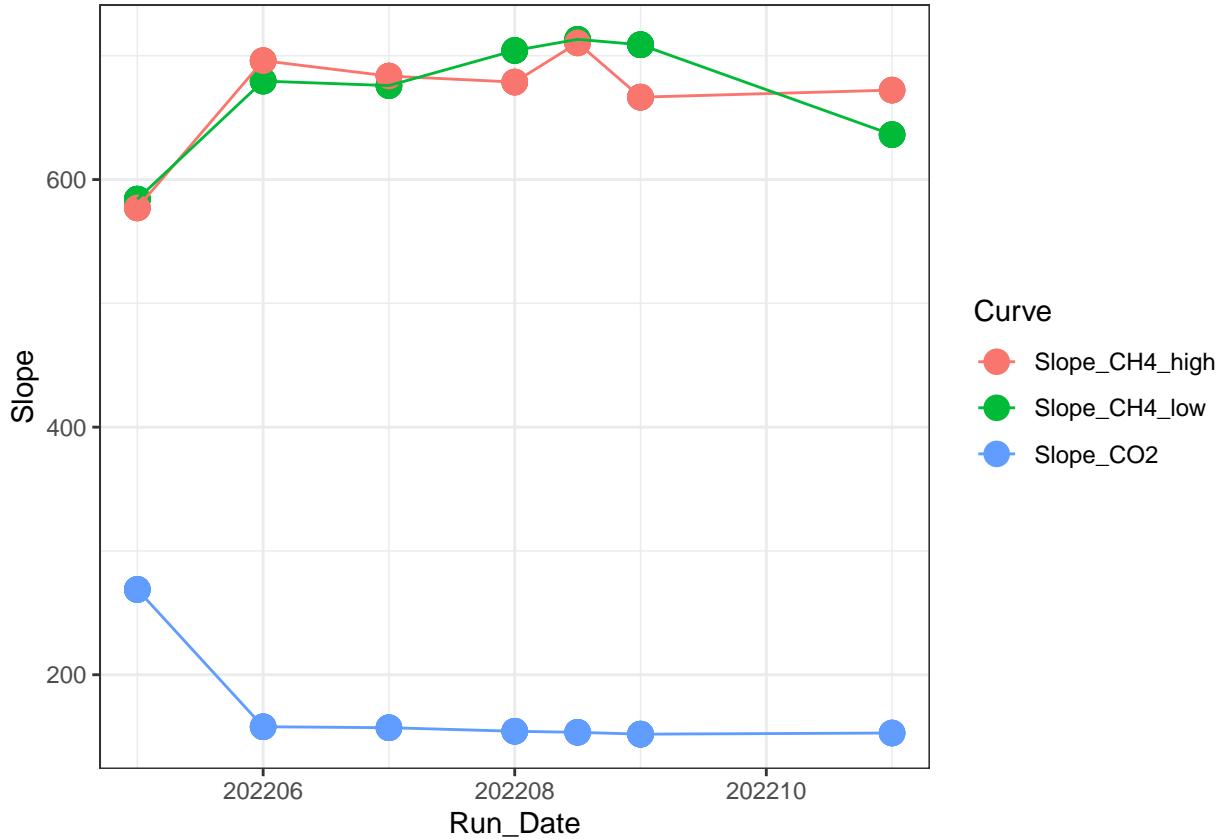
```
##  
## Call:  
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)  
##  
## Residuals:  
##      Min       1Q   Median       3Q      Max  
## -70767    1755    2118    3037   87949  
##  
## Coefficients:  
##                               Estimate Std. Error t value Pr(>|t|)  
## (Intercept)           -1829.347  11582.865  -0.158   0.877  
## stds_ch4$STD_Conc     577.146      5.454 105.826 <2e-16 ***  
## ---  
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1  
##  
## Residual standard error: 35760 on 12 degrees of freedom  
## Multiple R-squared:  0.9989, Adjusted R-squared:  0.9988  
## F-statistic: 1.12e+04 on 1 and 12 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:
##    Min     1Q Median     3Q    Max 
## -81062 -50011 -32599  16209 147447 
##
## Coefficients:
##             Estimate Std. Error t value Pr(>|t|)    
## (Intercept) 82323.28   44265.89   1.860 0.112263  
## stds_co2$STD_Conc 268.88      34.68   7.752 0.000242 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 99010 on 6 degrees of freedom
## Multiple R-squared:  0.9092, Adjusted R-squared:  0.8941 
## F-statistic: 60.1 on 1 and 6 DF,  p-value: 0.0002421

##      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)

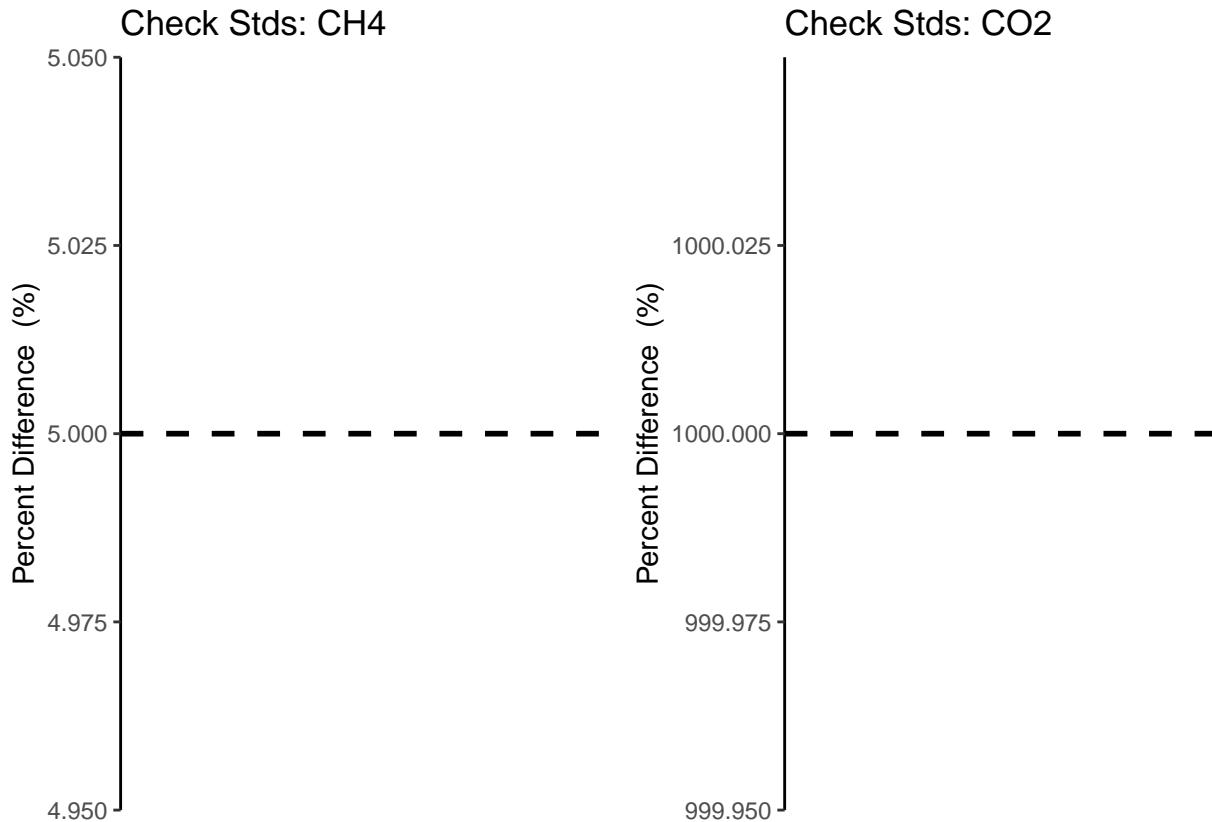
```

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)

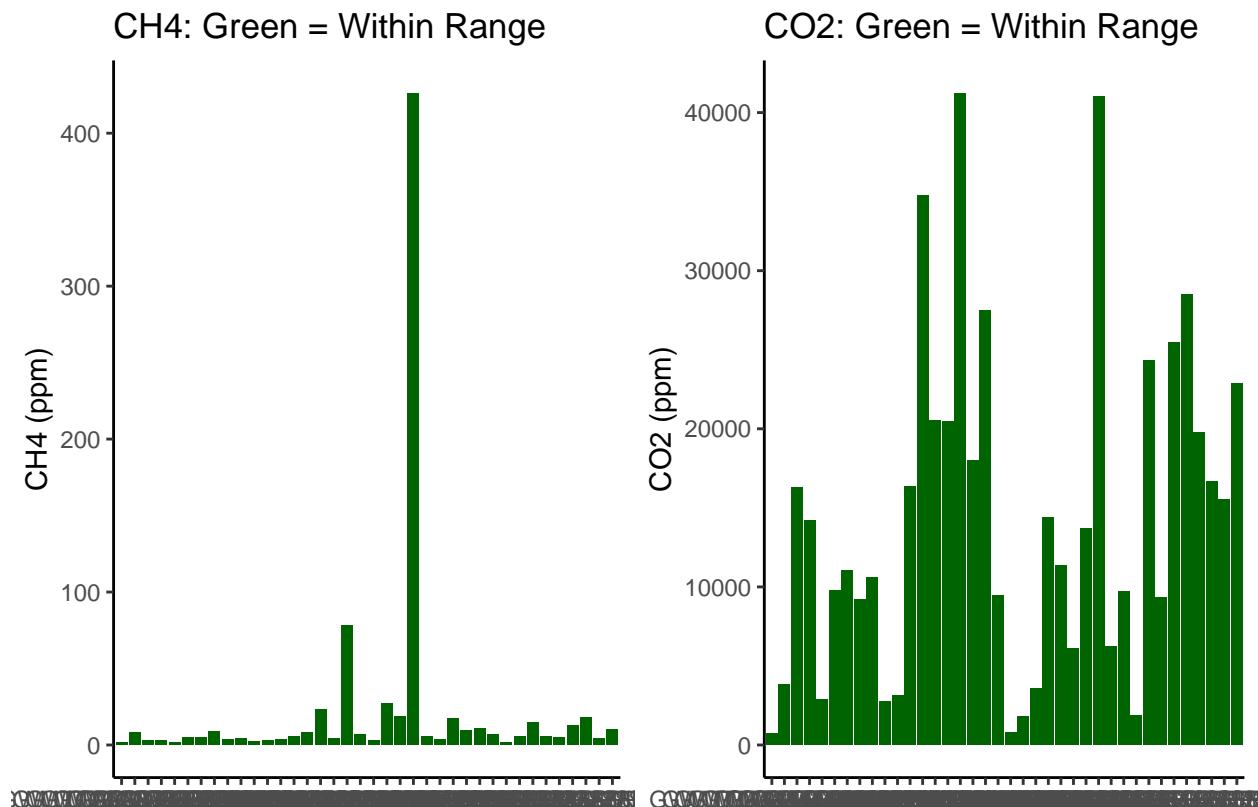
#check results
#head(Samples)

#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( "darkgreen", "red"))+
  #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 202205 Unknown  TGAS     2022
## 2 Varian GC Stephanie J. Wilson 202205 Unknown  TGAS     2022
## 3 Varian GC Stephanie J. Wilson 202205 Unknown  TGAS     2022
## 4 Varian GC Stephanie J. Wilson 202205 Unknown  TGAS     2022
## 5 Varian GC Stephanie J. Wilson 202205 Unknown  TGAS     2022
## 6 Varian GC Stephanie J. Wilson 202205 Unknown  TGAS     2022
##   Sample_Month    Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      May GWI_TGAS_TR_DS_1           2       NA    1420  285551
## 2      May GWI_TGAS_TR_DS_2           2       NA    5068 1114202
## 3      May GWI_TGAS_TR_DS_3           2       NA    2119 4462563
## 4      May GWI_TGAS_TR_SF_1           2       NA    1847 3899395
## 5      May GWI_TGAS_TR_SF_2           2       NA    1424  856806
## 6      May GWI_TGAS_TR_SF_3           2       NA    3065 2703534
##   Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm      CH4_Flag
## 1        NA        NA      Low    1.929804    755.8298 Within Range
## 2        NA        NA      Low    8.174561   3837.6884 Within Range
## 3        NA        NA      Low   3.126373  16290.6695 Within Range
## 4        NA        NA      Low   2.660755 14196.1760 Within Range
## 5        NA        NA      Low   1.936651 2880.3999 Within Range
## 6        NA        NA      Low   4.745765  9748.6166 Within Range
##   CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range            3.859607           1511.660
## 2 Within Range           16.349122           7675.377
## 3 Within Range            6.252746           32581.339
## 4 Within Range            5.321510           28392.352
## 5 Within Range            3.873302           5760.800
## 6 Within Range            9.491529           19497.233
```

```
#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 202205  TGAS     2022      May
## 2 Varian GC Stephanie J. Wilson 202205  TGAS     2022      May
## 3 Varian GC Stephanie J. Wilson 202205  TGAS     2022      May
## 4 Varian GC Stephanie J. Wilson 202205  TGAS     2022      May
## 5 Varian GC Stephanie J. Wilson 202205  TGAS     2022      May
## 6 Varian GC Stephanie J. Wilson 202205  TGAS     2022      May
##   Sample_ID Dilution_Factor Field.Notes      CH4_Flag      CO2_Flag
## 1 GWI_TGAS_TR_DS_1           2       NA Within Range Within Range
## 2 GWI_TGAS_TR_DS_2           2       NA Within Range Within Range
## 3 GWI_TGAS_TR_DS_3           2       NA Within Range Within Range
## 4 GWI_TGAS_TR_SF_1           2       NA Within Range Within Range
## 5 GWI_TGAS_TR_SF_2           2       NA Within Range Within Range
## 6 GWI_TGAS_TR_SF_3           2       NA Within Range Within Range
```

```

##    CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1          3.859607      1511.660
## 2         16.349122      7675.377
## 3         6.252746      32581.339
## 4         5.321510      28392.352
## 5         3.873302      5760.800
## 6         9.491529      19497.233

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 ,
         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_202205_Processed.csv")

#end

```