

COMPASS_Synoptic_TGW_2022: June

Stephanie J. Wilson

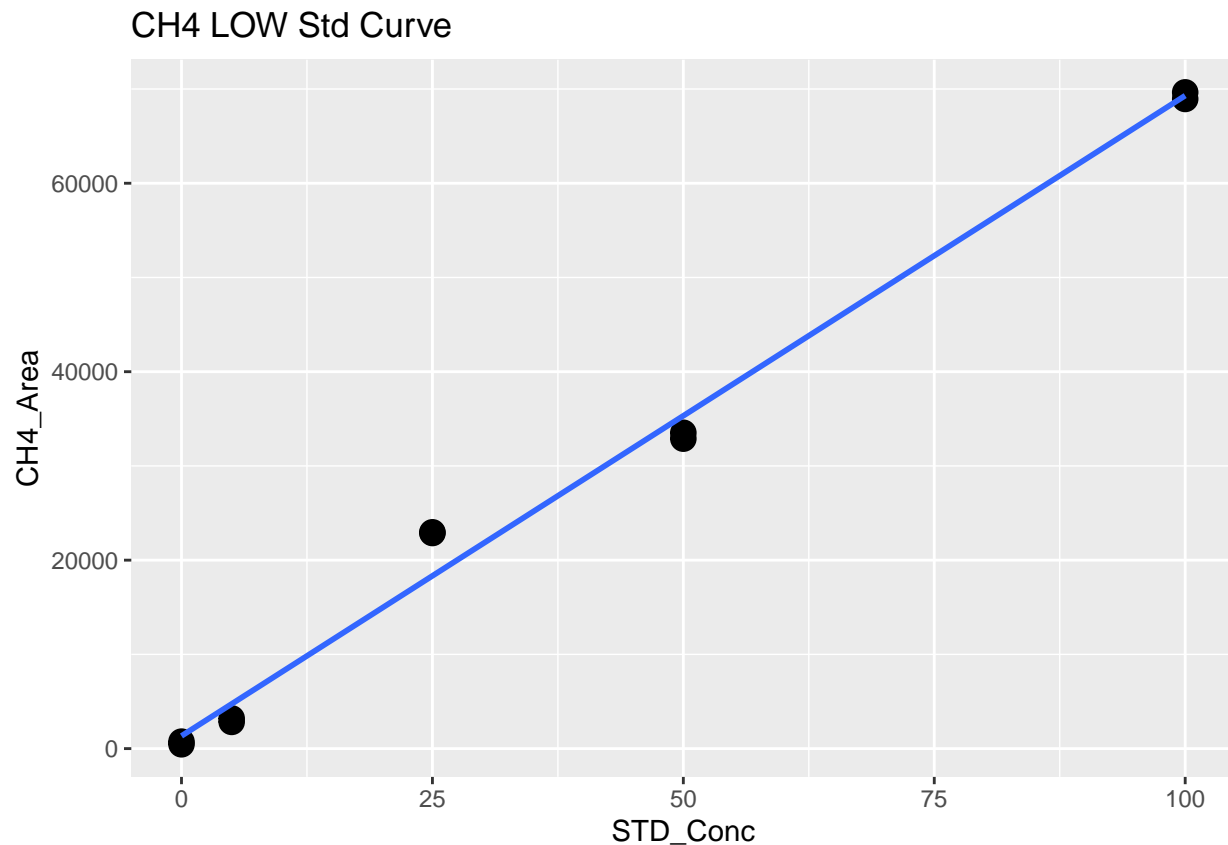
2023-01-14

##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 202206      Blank  TGAS      2022
## 2 Varian GC Stephanie J. Wilson 202206      Unknown TGAS      2022
## 3 Varian GC Stephanie J. Wilson 202206      Unknown TGAS      2022
## 4 Varian GC Stephanie J. Wilson 202206      Unknown TGAS      2022
## 5 Varian GC Stephanie J. Wilson 202206      Unknown TGAS      2022
## 6 Varian GC Stephanie J. Wilson 202206      Unknown TGAS      2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1      June      Blank_1      1      NA      0      0
## 2      June MSM_TGAS_UP_SF_1      2      NA      12697 1791814
## 3      June MSM_TGAS_UP_SF_2      2      NA      23232 5697789
## 4      June MSM_TGAS_UP_SF_3      2      NA      8442 10718524
## 5      June MSM_TGAS_UP_SF_4      2      NA      8731 6094426
## 6      June MSM_TGAS_UP_SF_5      2      NA      34611 8661859
##      Field.Notes Lab.Notes
## 1      NA      NA
## 2      NA      NA
## 3      NA      NA
## 4      NA      NA
## 5      NA      NA
## 6      NA      NA

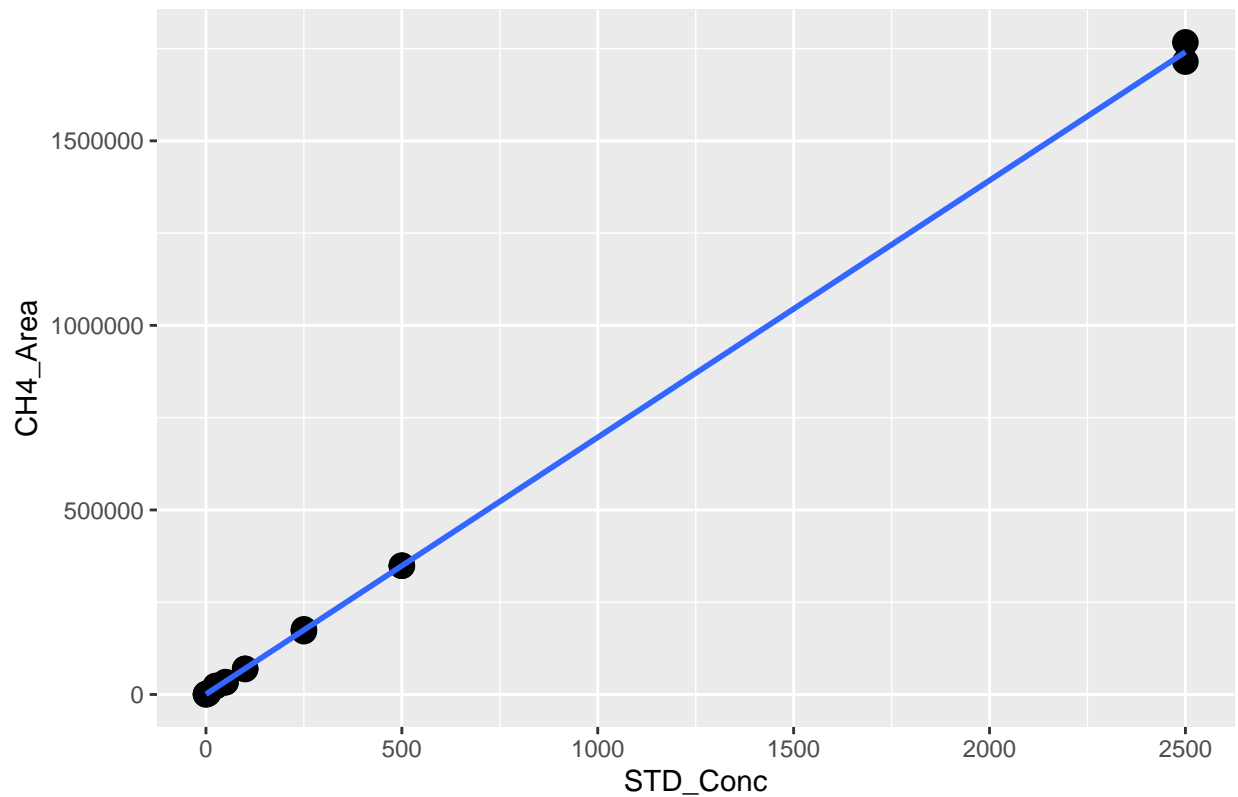
## 'geom_smooth()' using formula = 'y ~ x'
```



```
##
## Call:
## lm(formula = stds_ch4_low$CH4_Area ~ stds_ch4_low$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -2430.8 -1733.7  -744.1   173.6  4623.8
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    1342.56    1201.65   1.117   0.296
## stds_ch4_low$STD_Conc    679.55      23.43  29.001 2.16e-09 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 2706 on 8 degrees of freedom
## Multiple R-squared:  0.9906, Adjusted R-squared:  0.9894
## F-statistic: 841.1 on 1 and 8 DF, p-value: 2.163e-09

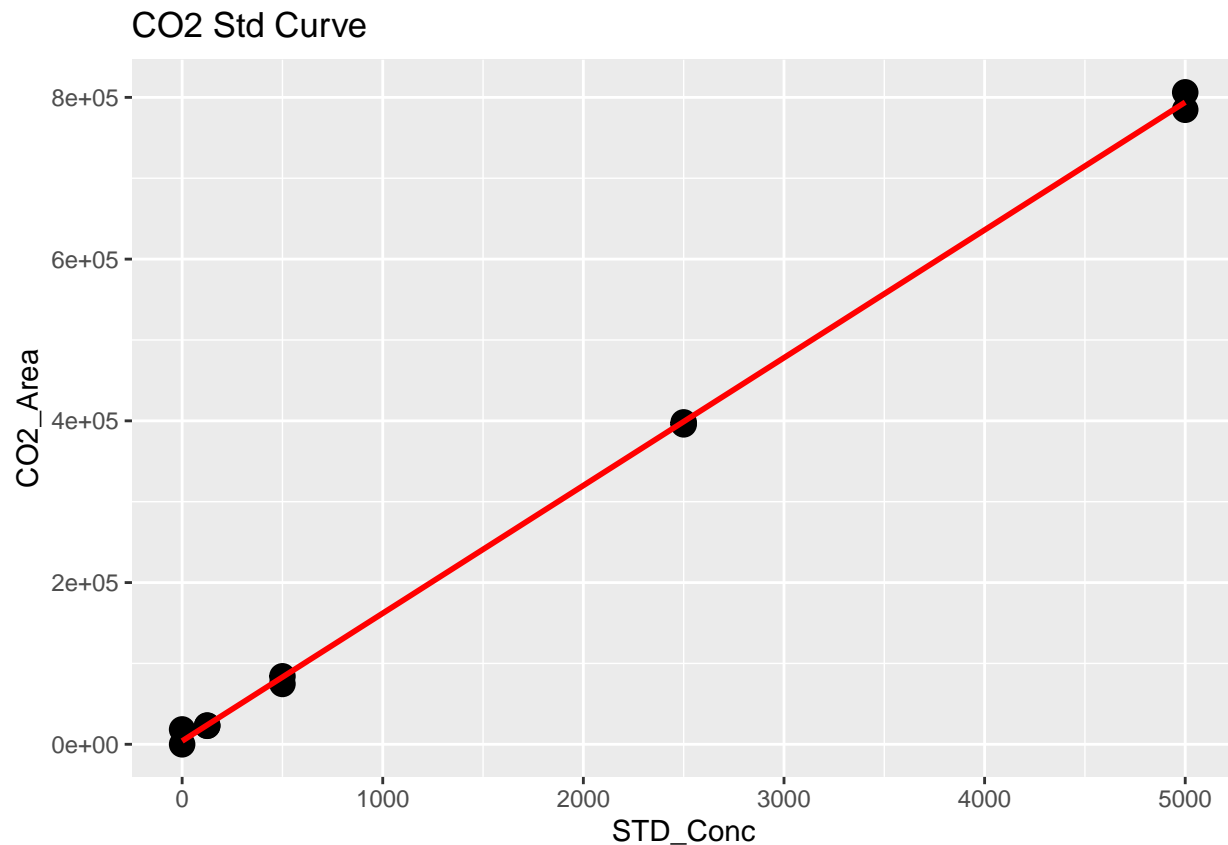
## 'geom_smooth()' using formula = 'y ~ x'
```

CH4 HIGH Std Curve



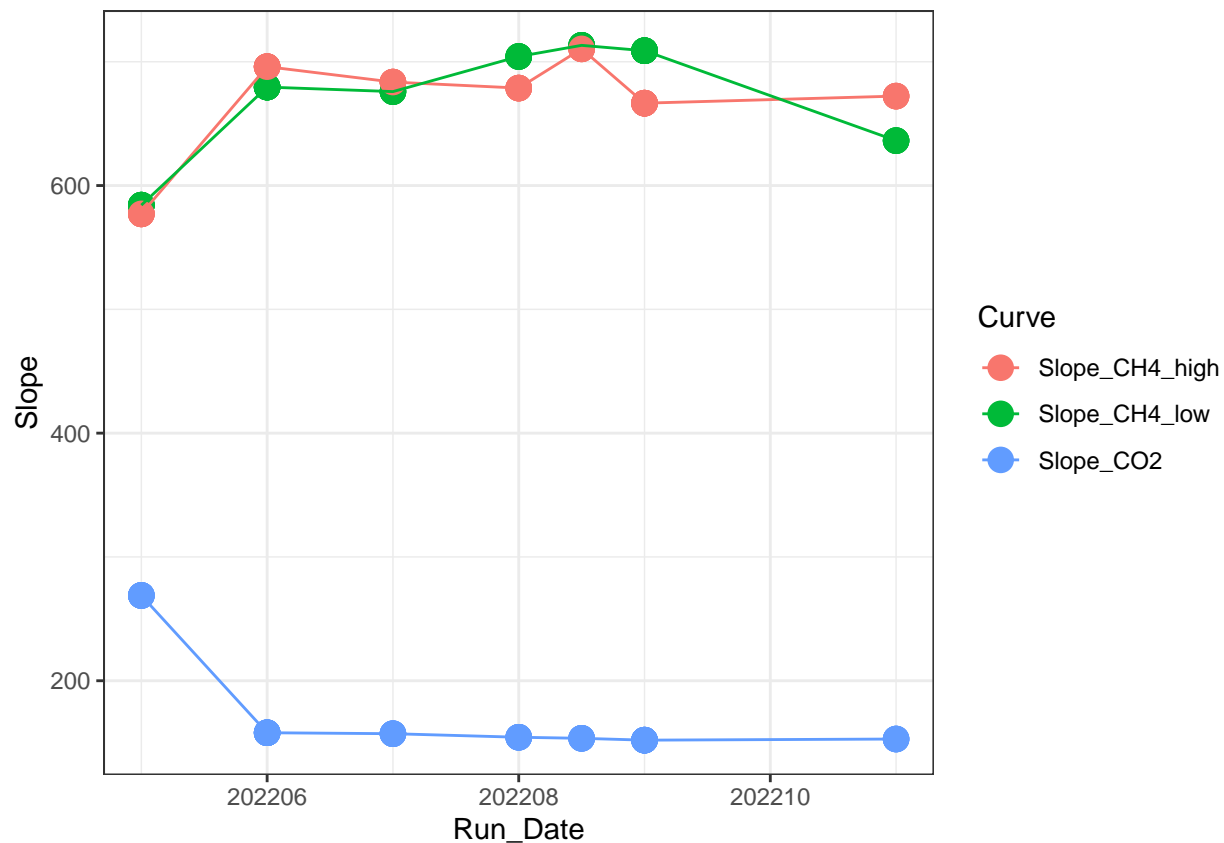
```
##
## Call:
## lm(formula = stds_ch4$CH4_Area ~ stds_ch4$STD_Conc)
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -26065.1  -1479.7   -414.3    670.4   26209.9
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)      664.824    2880.168   0.231   0.821
## stds_ch4$STD_Conc  696.019      3.177  219.093  <2e-16 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 10150 on 14 degrees of freedom
## Multiple R-squared:  0.9997, Adjusted R-squared:  0.9997
## F-statistic: 4.8e+04 on 1 and 14 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
## -9295.7 -3873.2  -928.8   611.6 14324.4
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)    4049.58    3415.57   1.186   0.27
## stds_co2$STD_Conc  157.99      1.36 116.130 3.38e-14 ***
## ---
## Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 8233 on 8 degrees of freedom
## Multiple R-squared:  0.9994, Adjusted R-squared:  0.9993
## F-statistic: 1.349e+04 on 1 and 8 DF, p-value: 3.379e-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)

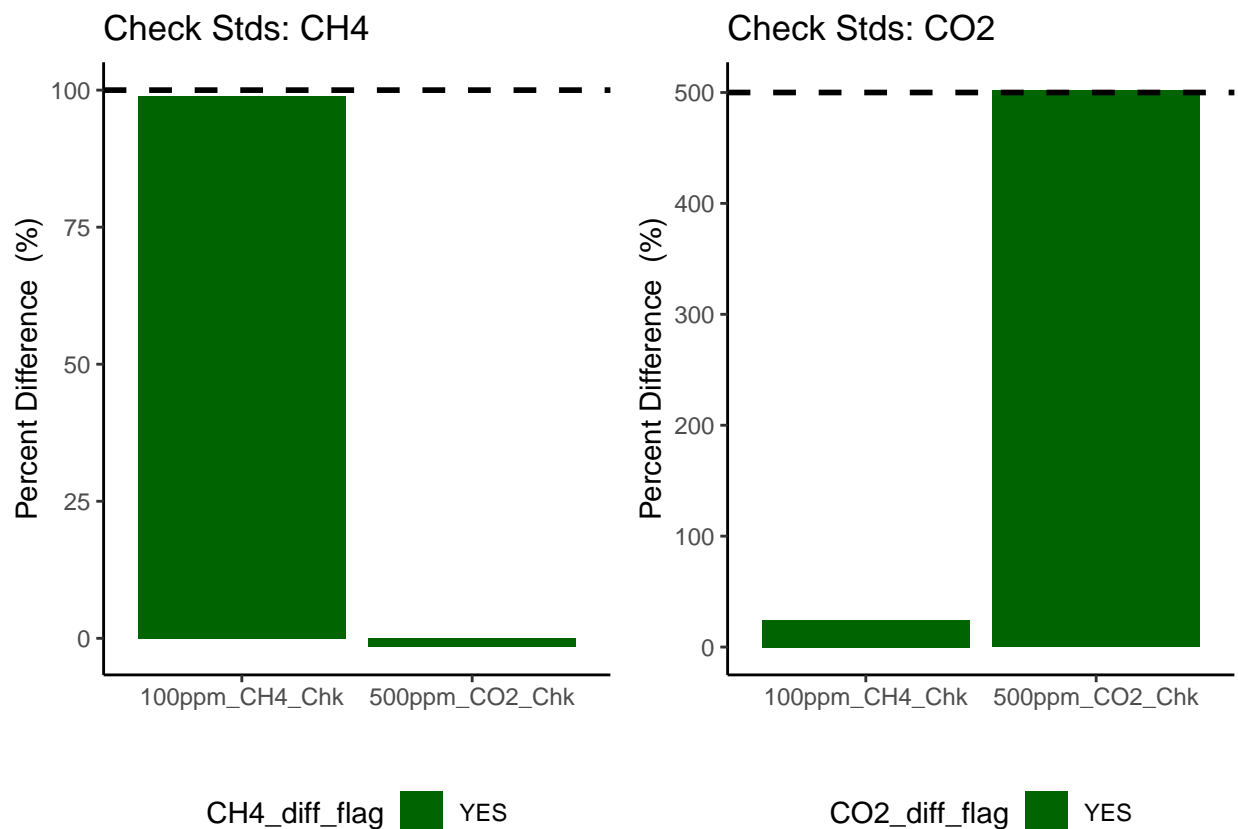
```

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.

```



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)

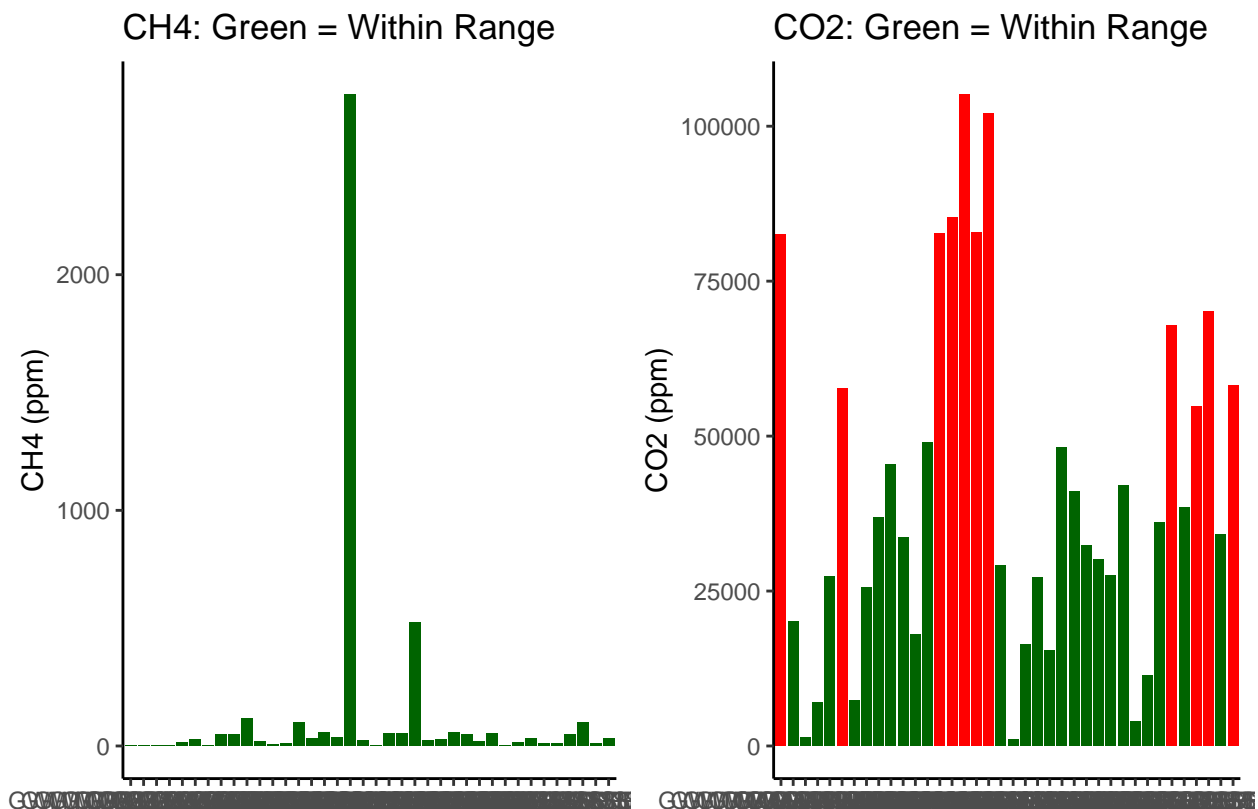
#check results
#head(Samples)

#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( "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)

```



Write out processed data & slopes

```
#check results
head(Samples)
```

```
##      Machine      User Run_Date Sample_Type Type1 Sample_Year
## 1 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
## 2 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
## 3 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
## 4 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
## 5 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
## 6 Varian GC Stephanie J. Wilson 202206 Unknown TGAS 2022
##      Sample_Month      Sample_ID Dilution_Factor STD_Conc CH4_Area CO2_Area
## 1 June MSM_TGAS_UP_SF_1 2 NA 12697 1791814
## 2 June MSM_TGAS_UP_SF_2 2 NA 23232 5697789
## 3 June MSM_TGAS_UP_SF_3 2 NA 8442 10718524
## 4 June MSM_TGAS_UP_SF_4 2 NA 8731 6094426
## 5 June MSM_TGAS_UP_SF_5 2 NA 34611 8661859
## 6 June MSM_TGAS_UP_SF_6 2 NA 68050 11080250
##      Field.Notes Lab.Notes CH4_Curve CH4_Conc_ppm CO2_Conc_ppm CH4_Flag
## 1 NA NA Low 16.70887 11315.66 Within Range
## 2 NA NA Low 32.21188 36038.55 Within Range
## 3 NA NA Low 10.44734 67817.32 Within Range
## 4 NA NA Low 10.87262 38549.07 Within Range
## 5 NA NA Low 48.95689 54799.65 Within Range
## 6 NA NA Low 98.16478 70106.87 Within Range
##      CO2_Flag CH4_Conc_ppm_dilcorr CO2_Conc_ppm_dilcorr
## 1 Within Range 33.41775 22631.33
## 2 Within Range 64.42376 72077.11
## 3 Needs Dilution 20.89467 135634.65
## 4 Within Range 21.74524 77098.14
## 5 Needs Dilution 97.91379 109599.30
## 6 Needs Dilution 196.32956 140213.74
```

```
#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 202206 TGAS 2022 June
## 2 Varian GC Stephanie J. Wilson 202206 TGAS 2022 June
## 3 Varian GC Stephanie J. Wilson 202206 TGAS 2022 June
## 4 Varian GC Stephanie J. Wilson 202206 TGAS 2022 June
## 5 Varian GC Stephanie J. Wilson 202206 TGAS 2022 June
## 6 Varian GC Stephanie J. Wilson 202206 TGAS 2022 June
##      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 Needs Dilution
## 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          33.41775          22631.33
## 2          64.42376          72077.11
## 3          20.89467         135634.65
## 4          21.74524          77098.14
## 5          97.91379         109599.30
## 6         196.32956         140213.74
```

```
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_202206_Processed.csv")
```

```
#end
```