

# TEMPEST\_FW\_Well\_TSS\_20251110

Stephanie J. Wilson

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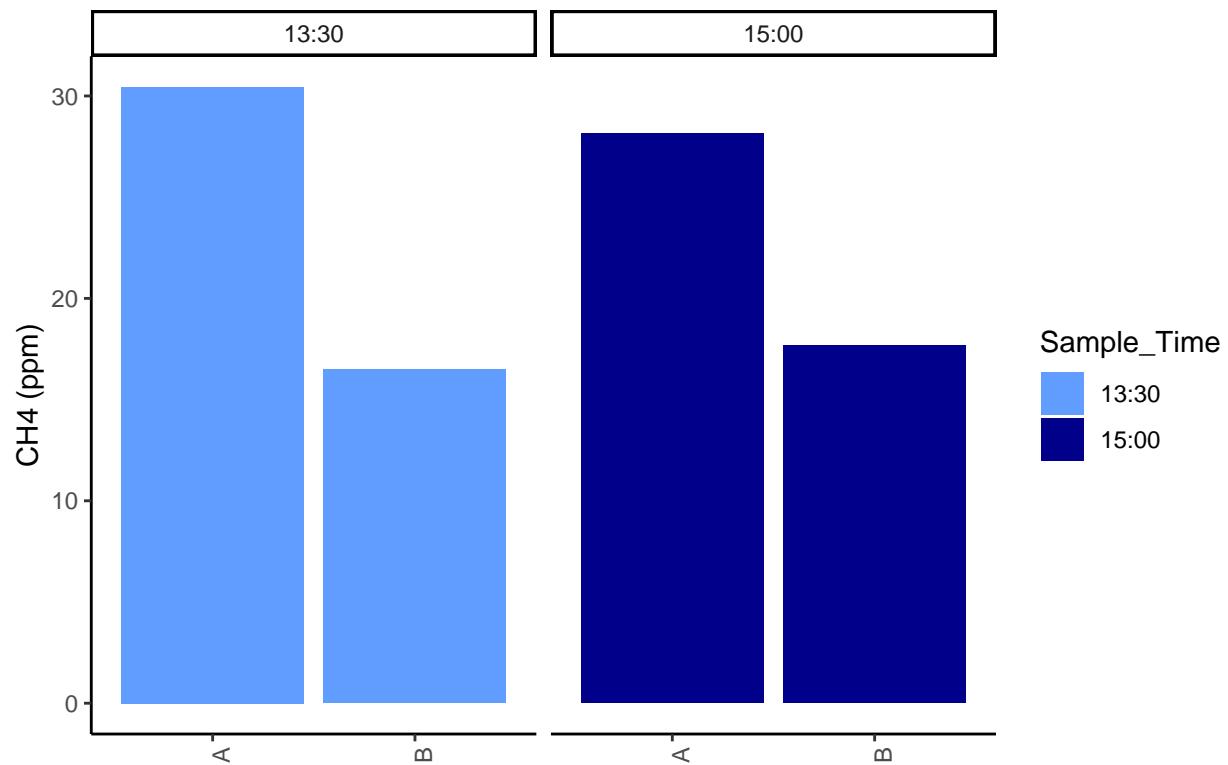
## Contents

0.1	GHG Data	2
0.2	DIC Data	3
0.3	TOC/TN Data	3
0.4	FE Data	5
0.5	Sulfate / Chloride Data	8
0.6	TSS Data	8

## 0.1 GHG Data

## Greenhouse Gas Data from Shimadzu GC - Shakey Samples

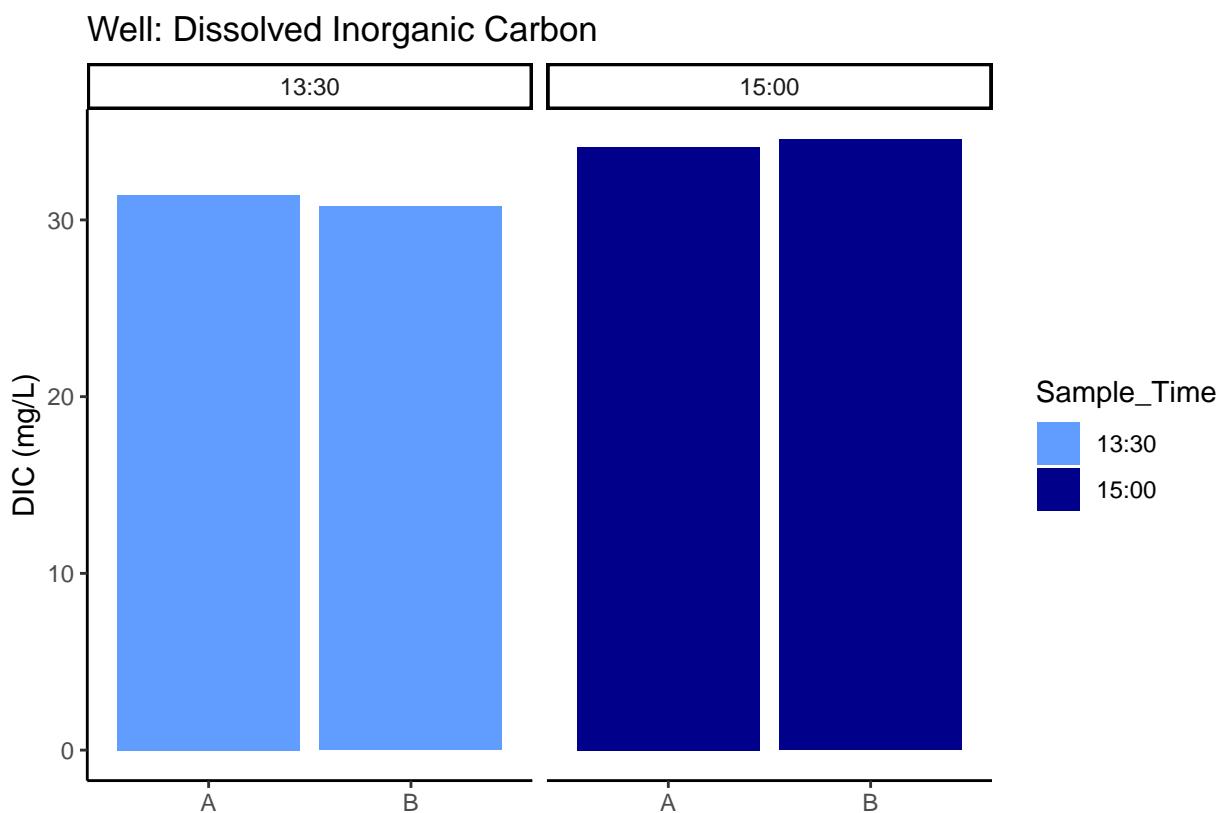
Well: Extractable Methane



Analyte	Average	Std_Deviation
CH4 (ppm)	23.18	7.14

## 0.2 DIC Data

```
## DIC Data from Shimadzu TOC-V
```

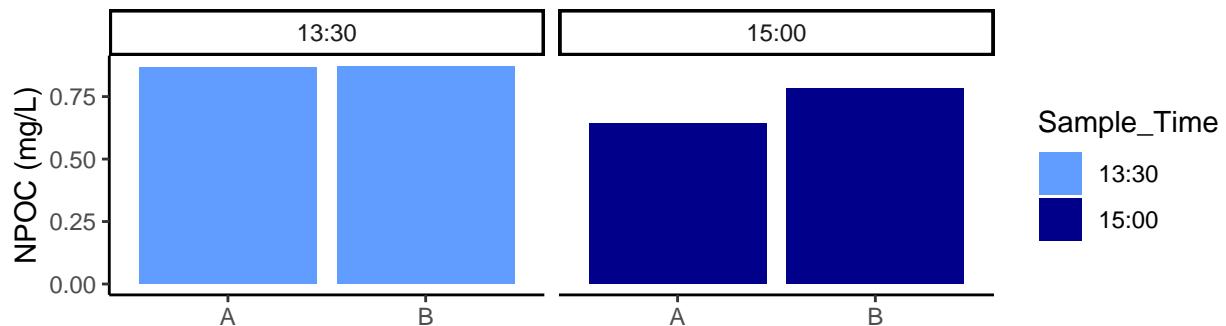


Analyte	Average	Std_Deviation
DIC (mg/L)	32.71	1.89

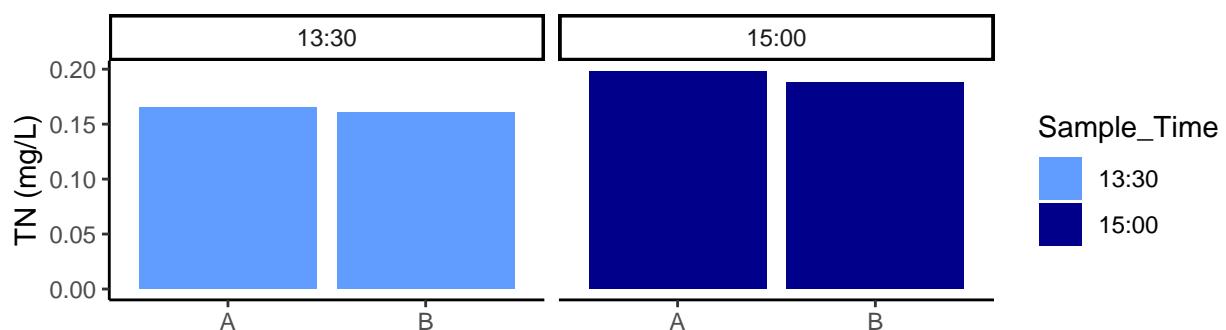
## 0.3 TOC/TN Data

```
## DOC and TN Data from Shimadzu TOC/TN
```

### Well: Dissolved Organic Carbon



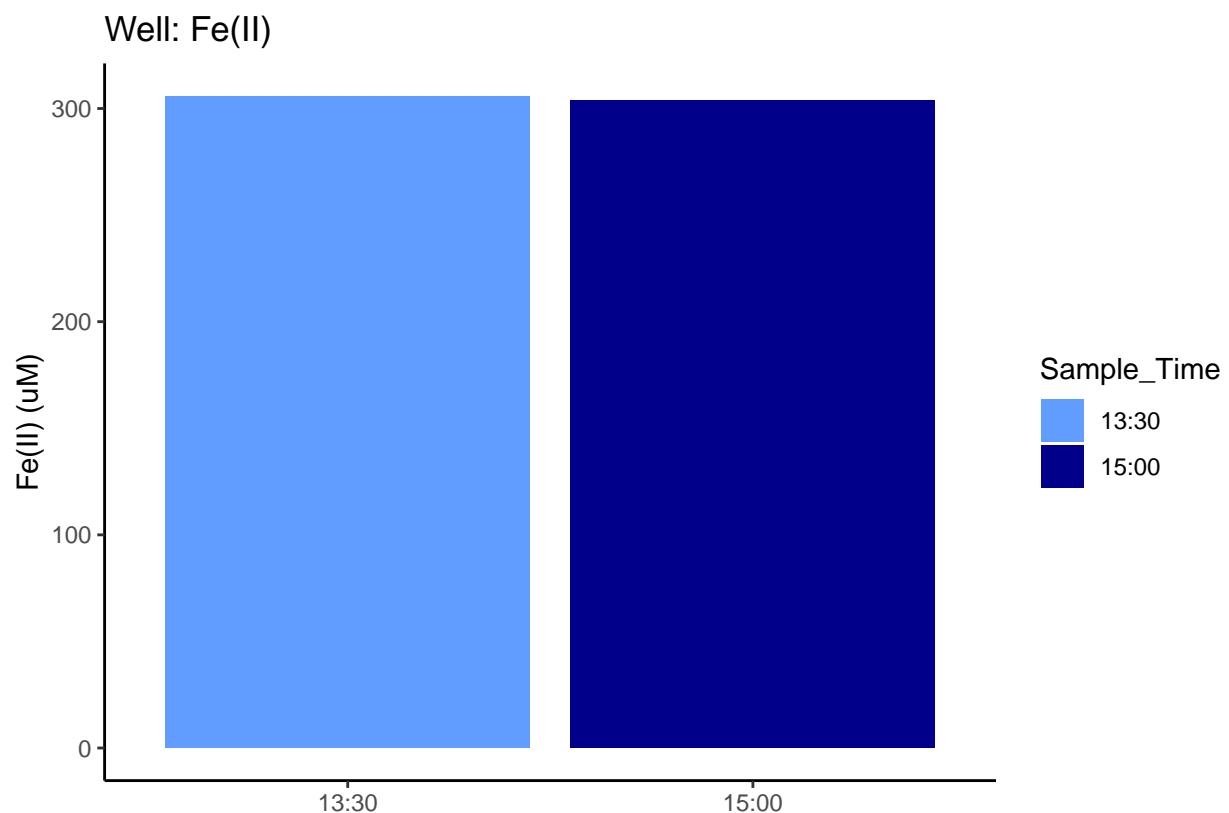
### Well: Total Nitrogen

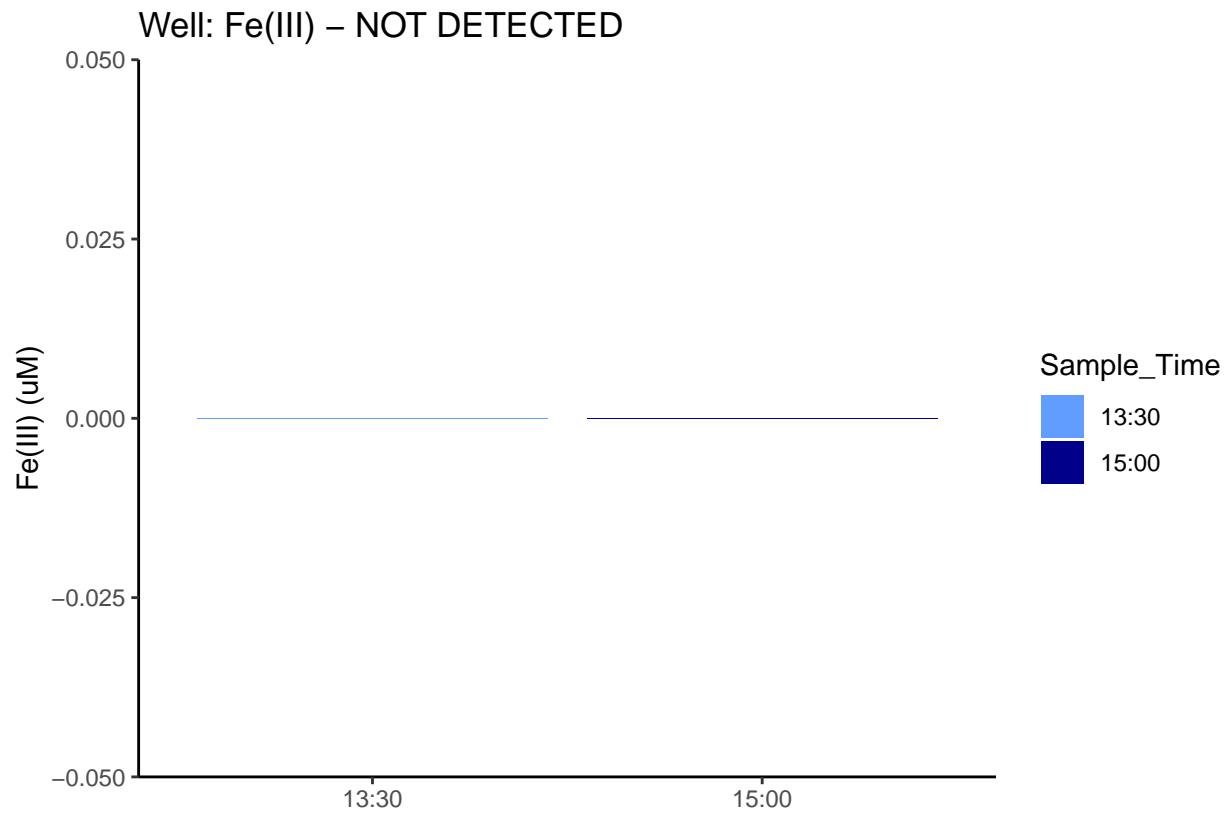


Analyte	Average	Std_Deviation
DOC (mg/L)	0.790	0.106
TN (mg/L)	0.178	0.018

## 0.4 FE Data

```
## FE Data from Shimadzu TOC-V
```

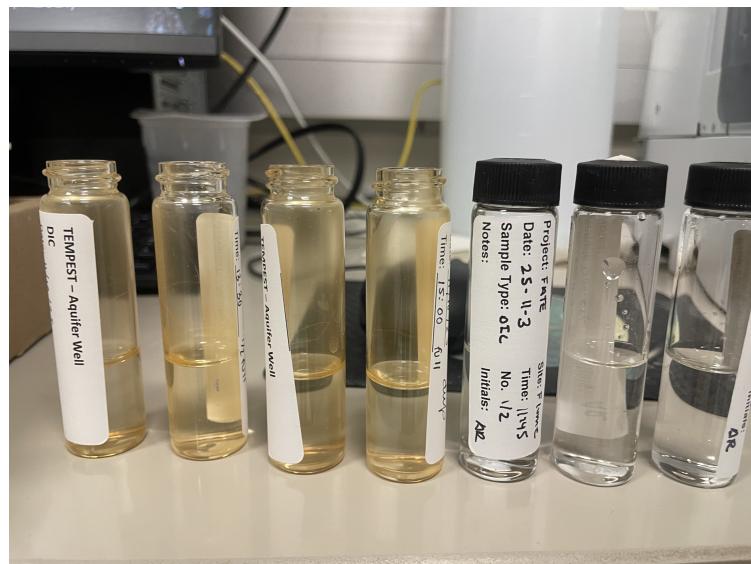




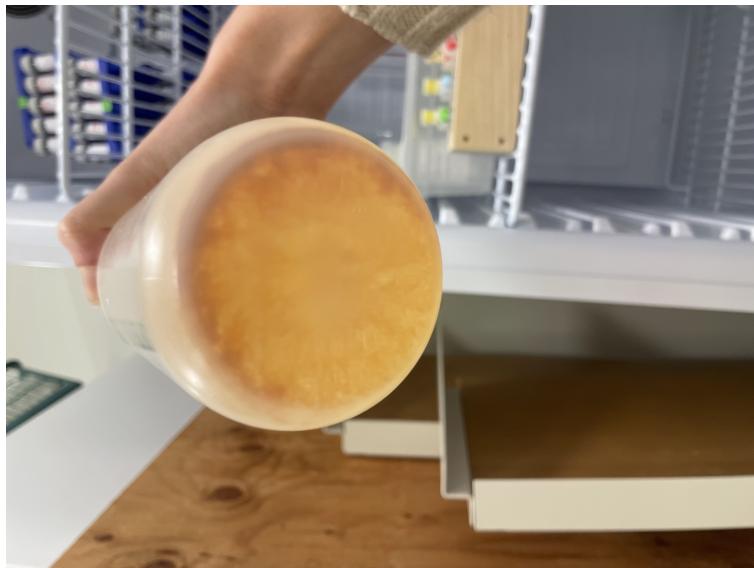
Analyte	Average	Std_Deviation
Fe(II) ( $\mu\text{M}$ )	304.9	1.36
Fe(III) ( $\mu\text{M}$ )	0.0	0.00

## Pictures of FE precipitation

## Four TMP Aquifer Well Vials compared to those with surface water from Rhode River:



## Refrigerated samples from TMP Aquifer Well:



#STILL WORKING ON THE REST OF THE SAMPLES:

## 0.5 Sulfate / Chloride Data

## 0.6 TSS Data

```
``{#r, echo=FALSE}

#read out title cat("TSS Data: ~1L Samples filtered with GFF")

#Read in the data tss <- read.csv("Raw Data/COMPASS_TEMPEST_WellTest_TSS_Data.csv")
#head(tss)

#Calculate an Avg. Sample Weight and caluclate TSS mg/L tss <- tss %>% mutate(Avg_Sample_Weight_g = rowMeans(select(., Tin_Filter_Sample_Weight_Dry_g, Tin_Filter_Sample_Weight_Dry1_g), na.rm = TRUE))

tssTSSmgL <- ((tssAvg_Sample_Weight_g - tssTinFilterWeightg)*1000)/(tssFiltered_Volume_mL/1000)

final_tss <- tss %>% #select(Project, Plot, grid, sample_name, Vial_ID, date, ) %>% mutate( Project = "COMPASS", # new column with same value on every row Experiment = "TEMPEST: Well Test", Sample_Date = "2025-09-04", Sample_Time = c("12:25", "13:00", "14:00", "NA"), Replicate = c("A", "A", "A", "NA"))

#this needs altered to match the tempest metadata and clean up final_tss <- final_tss %>% rename(sample_name = Sample_ID) %>% select( Project, Experiment, Sample_Date, Sample_Time, Replicate, sample_name, TSS_mgL, Filtered_Volume_mL # list columns in the order you want them )

#plot the TSS tss_plot <- ggplot(final_tss, aes(x=sample_name, y=TSS_mgL, fill=Sample_Time)) + geom_bar(stat = "identity") + labs(title="Well: Total Dissolved Solids", y="Total Dissolved Solids (mg/L)", x="") + theme_classic()

print(tss_plot)

##remove the DI blank and make a table tss_table <- final_tss %>% filter(sample_name != "DI_Bank") %>% select("sample_name", "TSS_mgL") %>% mutate(TSS_mgL = round(TSS_mgL, 2))

kable(tss_table)

##remove the DI blank and make a table final_tss <- final_tss %>% filter(sample_name != "DI_Bank")

#will put final data in processed data folder write.csv(final_tss, "Processed Data/TMP_20250904_FW_WellTest_TSS_Proc

````
```