

# Stream temperatures

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```
library(tidyverse)
library(lubridate)
# knitr::opts_chunk$set(fig.height = 3, fig.width = 6)
```

Explore stream temperatures for all sites.

*This is an example of where we might end up*

## Functions

```
# Calculate useful data coverage summaries
coverage_summary <- function(data){
  data %>%
  summarise(
    n_obs = n(),
    n_days = n_distinct(date),
    n_missing = sum(is.na(watertemp_mean_day))
  )
}
```

```
# Plot watertemp_day time series
# Note this is the two argument version
plot_temperature <- function(data, title){
  data %>%
  ggplot(aes(date, watertemp_mean_day)) +
  geom_line(aes(color = watertemp_method)) +
  scale_color_brewer("Method", palette = "Set2") +
  labs(x = "", y = "Water temperature (C)",
       title = title)
}
```

## Data Import

```
streams <- read_csv("data/HT00441_v8.csv")
names(streams) <- tolower(names(streams))
```

Filter to time period of interest, and set observations with anything other than “Accepted” quality to missing:

```
streams <- streams %>%
  filter(year(date) > 2000, year(date) < 2020) %>%
  mutate(
    watertemp_mean_day = ifelse(watertemp_mean_flag != "A",
      NA, watertemp_mean_day)) %>%
  select(sitecode, date, watertemp_mean_day, watertemp_method)
```

## Data coverage

coverage\_summary() will work on grouped data:

```
streams %>%
  group_by(sitecode) %>%
  coverage_summary()
```

## 'summarise()' ungrouping output (override with '.groups' argument)

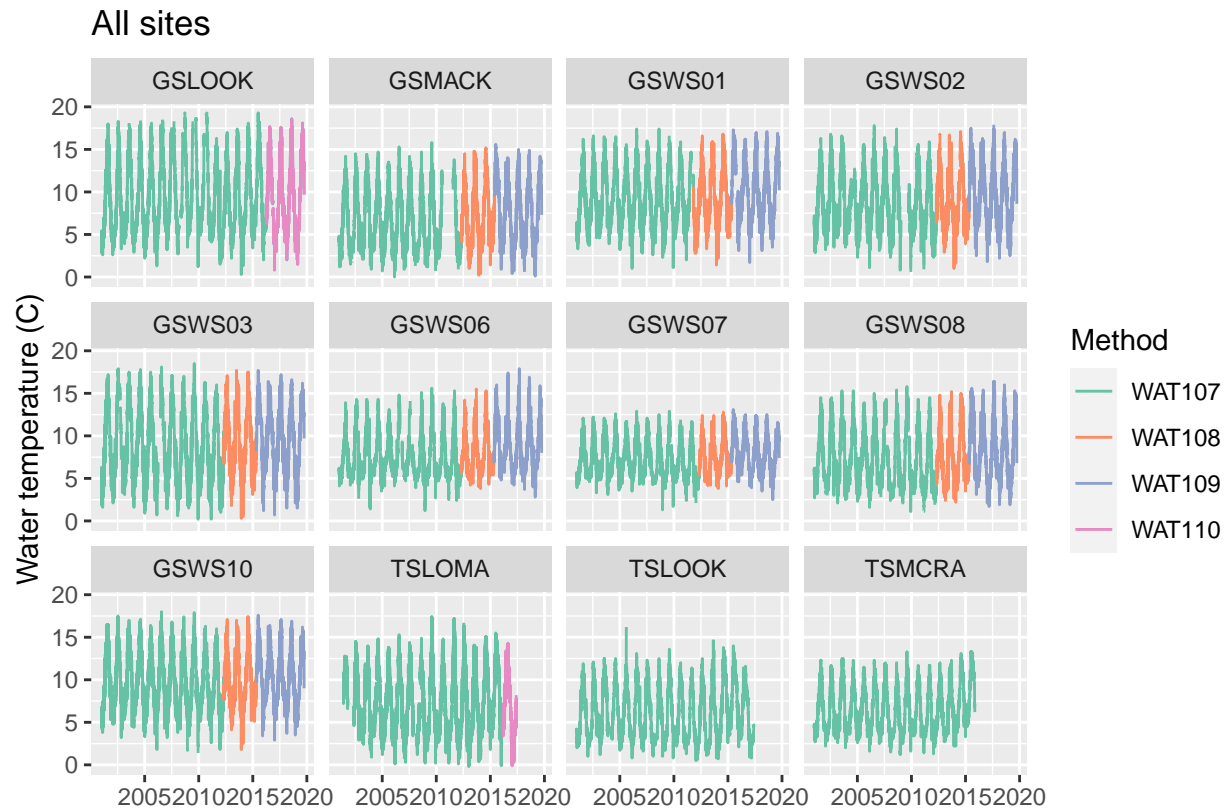
```
## # A tibble: 12 x 4
##   sitecode n_obs n_days n_missing
##   <chr>    <int> <int>    <int>
## 1 GSLOOK    8120   6841     543
## 2 GSMACK    6847   6847     428
## 3 GSWS01    6847   6847     223
## 4 GSWS02    6847   6847     566
## 5 GSWS03    6847   6847     179
## 6 GSWS06    6847   6847     432
## 7 GSWS07    6847   6847      73
## 8 GSWS08    6847   6847     236
## 9 GSWS10    6847   6847      85
## 10 TSLOMA   11976   5988     938
## 11 TSLOOK    5989   5989     115
## 12 TSMCRA    5427   5427      0
```

## Plots

One option is to apply the plot to all the data, then apply facetting to the resulting plot:

```
streams %>%
  plot_temperature("All sites") +
  facet_wrap(~ sitecode)
```

## Warning: Removed 10 row(s) containing missing values (geom\_path).



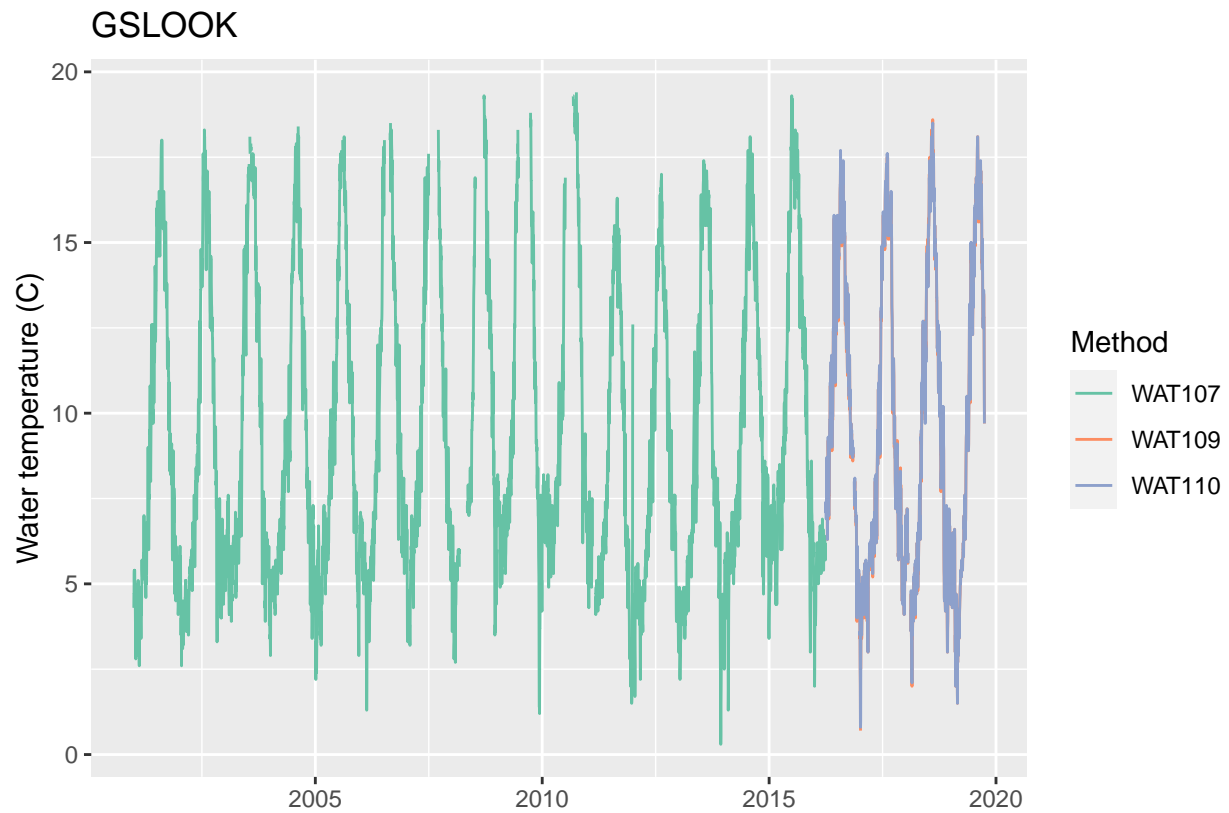
Alternatively, create and store all the plots with `rowwise()`:

```
streams_with_plots <- streams %>%
  nest_by(sitecode) %>%
  rowwise() %>%
  mutate(plots = list(plot_temperature(data, sitecode)))
```

Display one plot:

```
streams_with_plots$plots[[1]]
```

```
## Warning: Removed 5 row(s) containing missing values (geom_path).
```



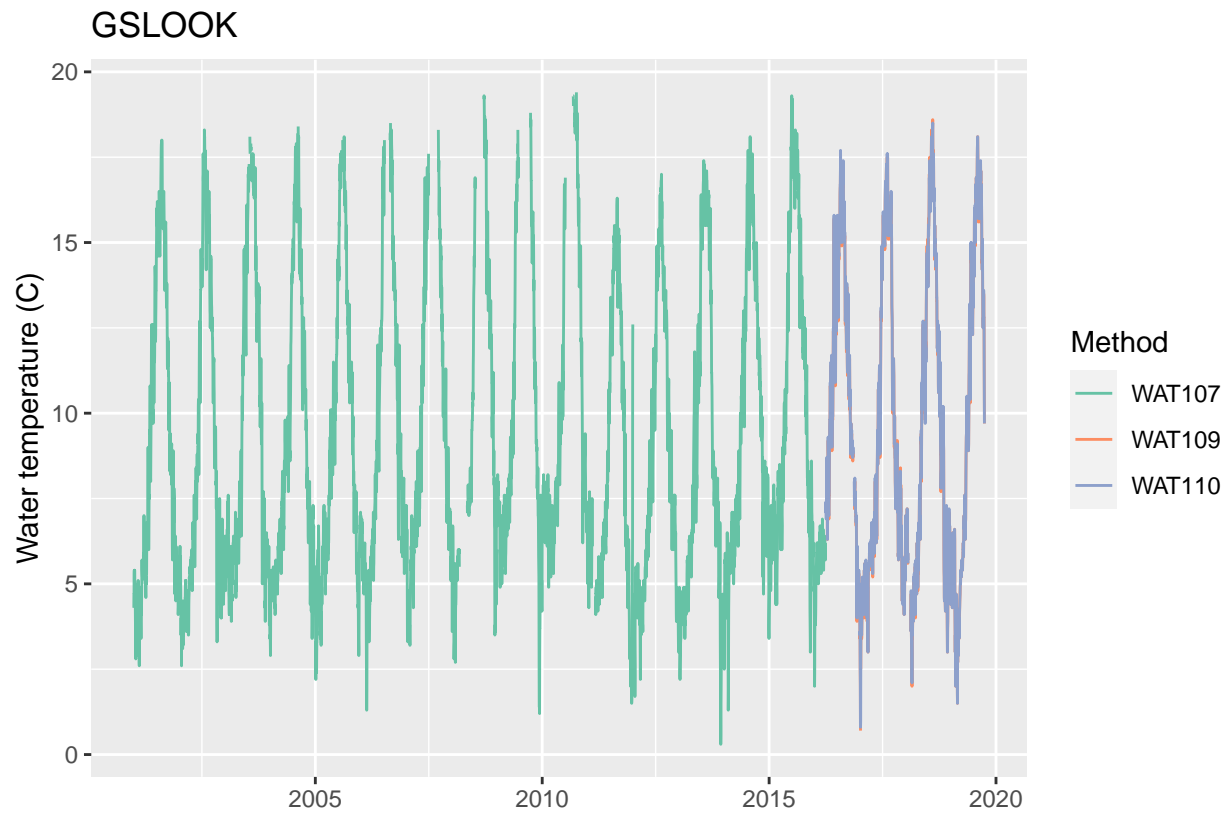
## Different ways to actually see all the plots

(Also a survey of other iteration methods)

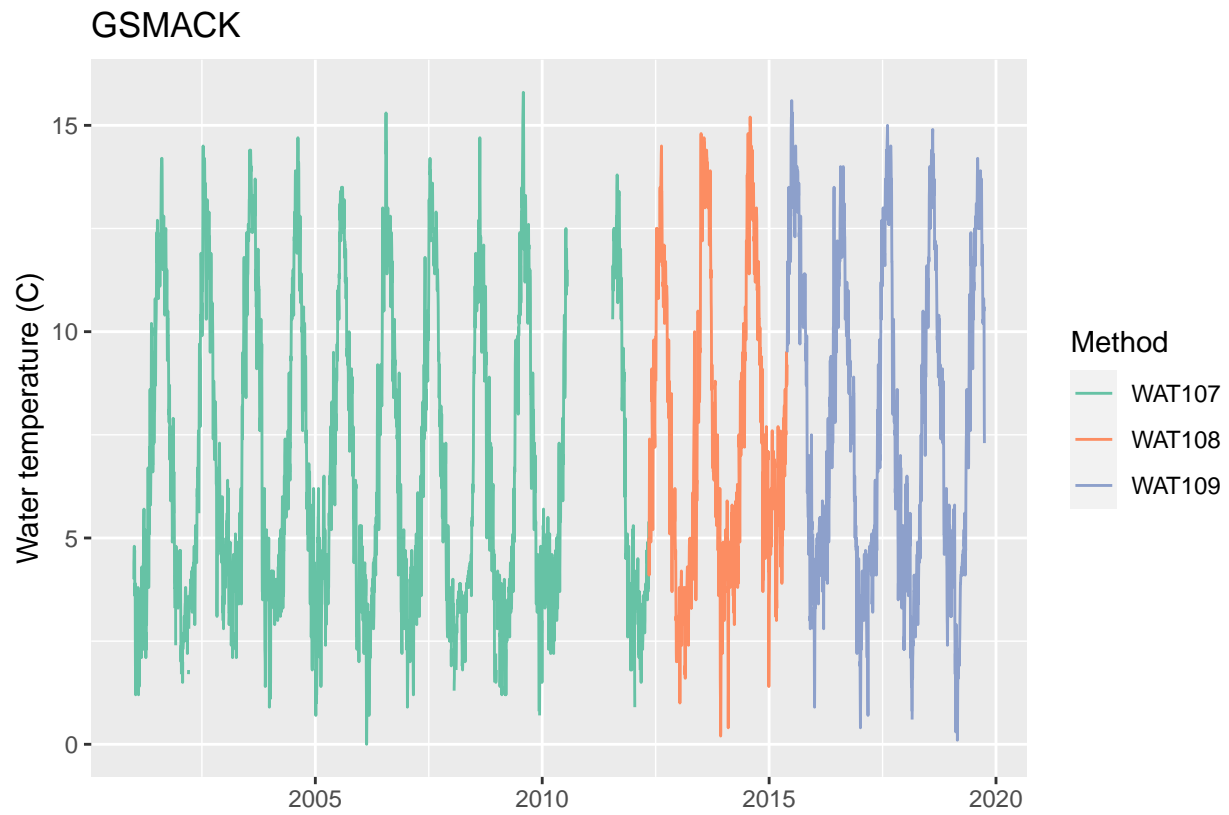
purrrs walk() function:

```
streams_with_plots %>% pull(plots) %>% walk(print)
```

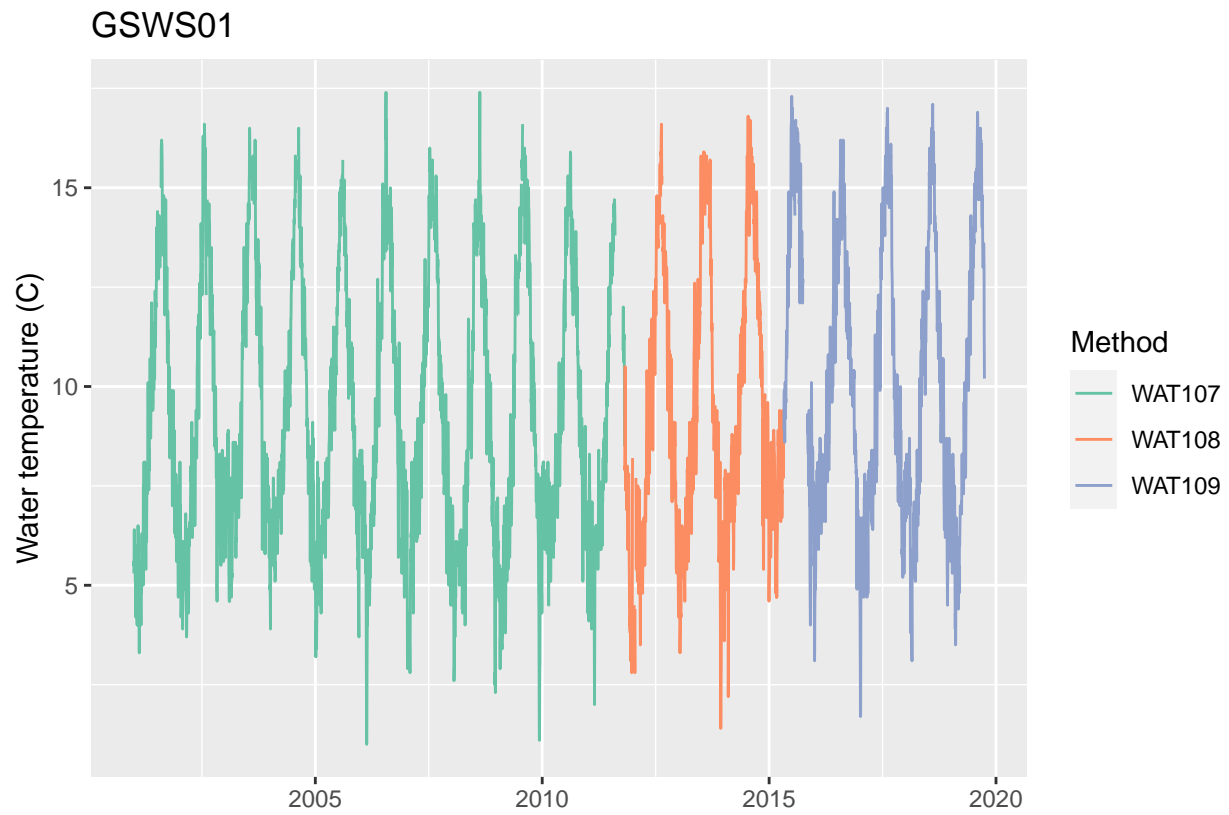
```
## Warning: Removed 5 row(s) containing missing values (geom_path).
```



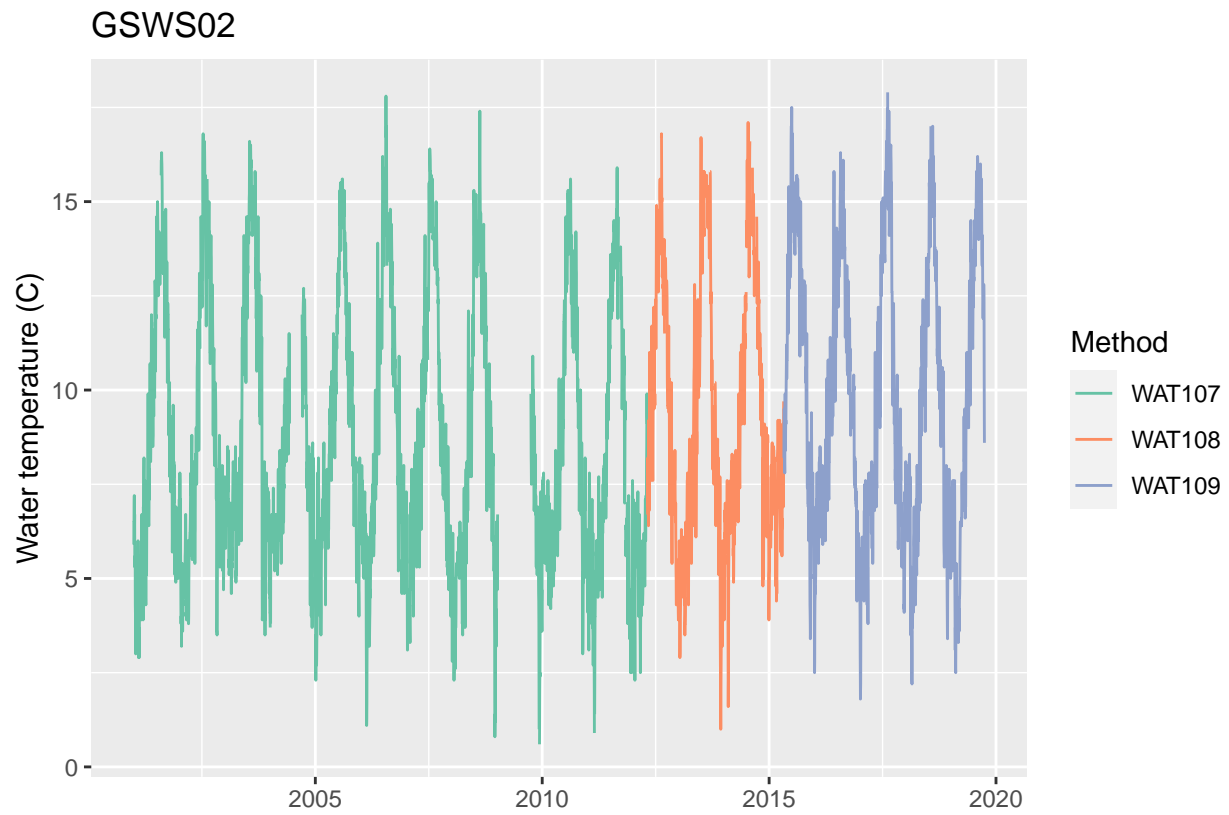
```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```



```
## Warning: Removed 2 row(s) containing missing values (geom_path).
```

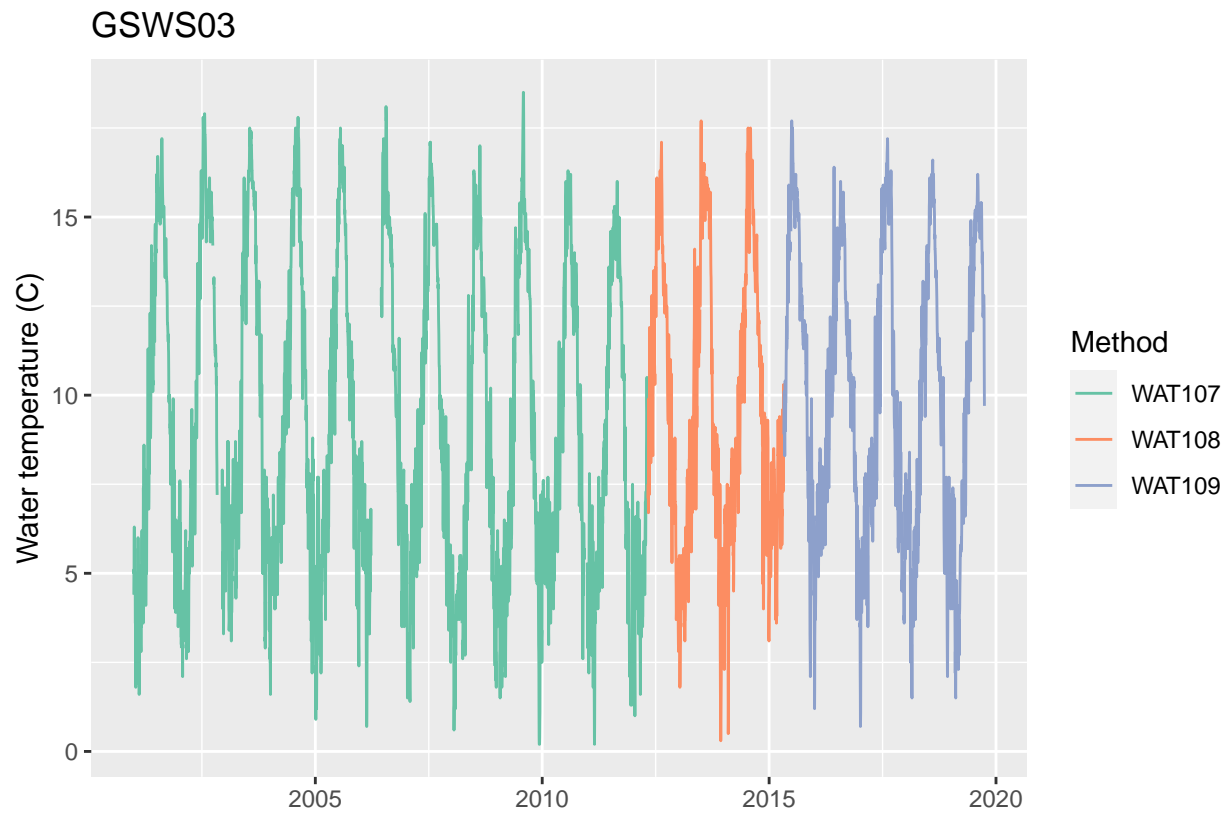


```
## Warning: Removed 2 row(s) containing missing values (geom_path).
```

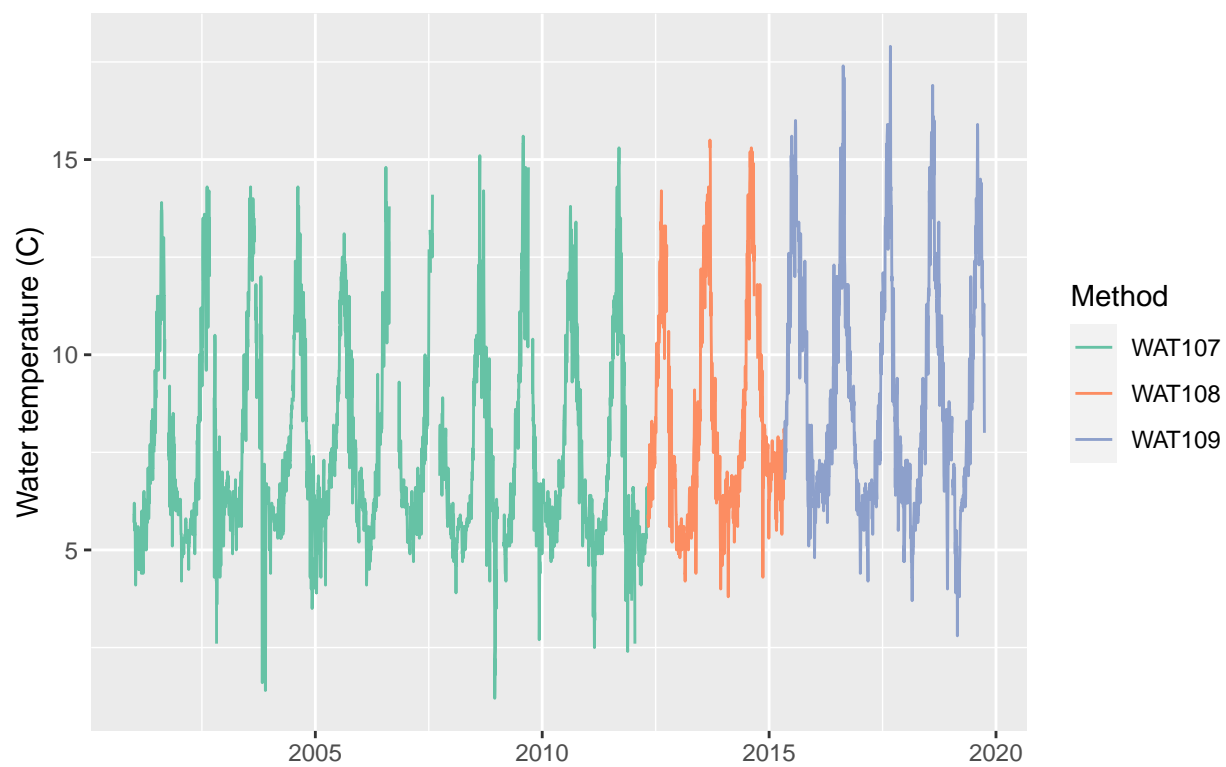


```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

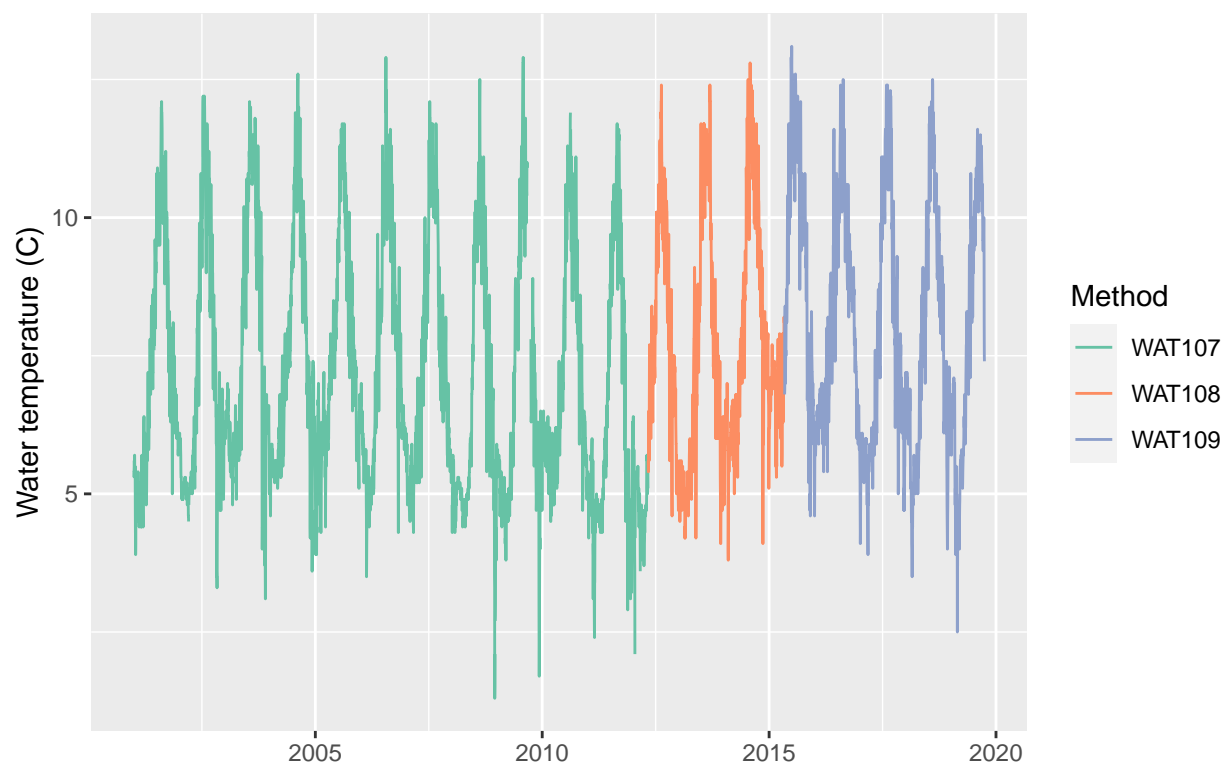


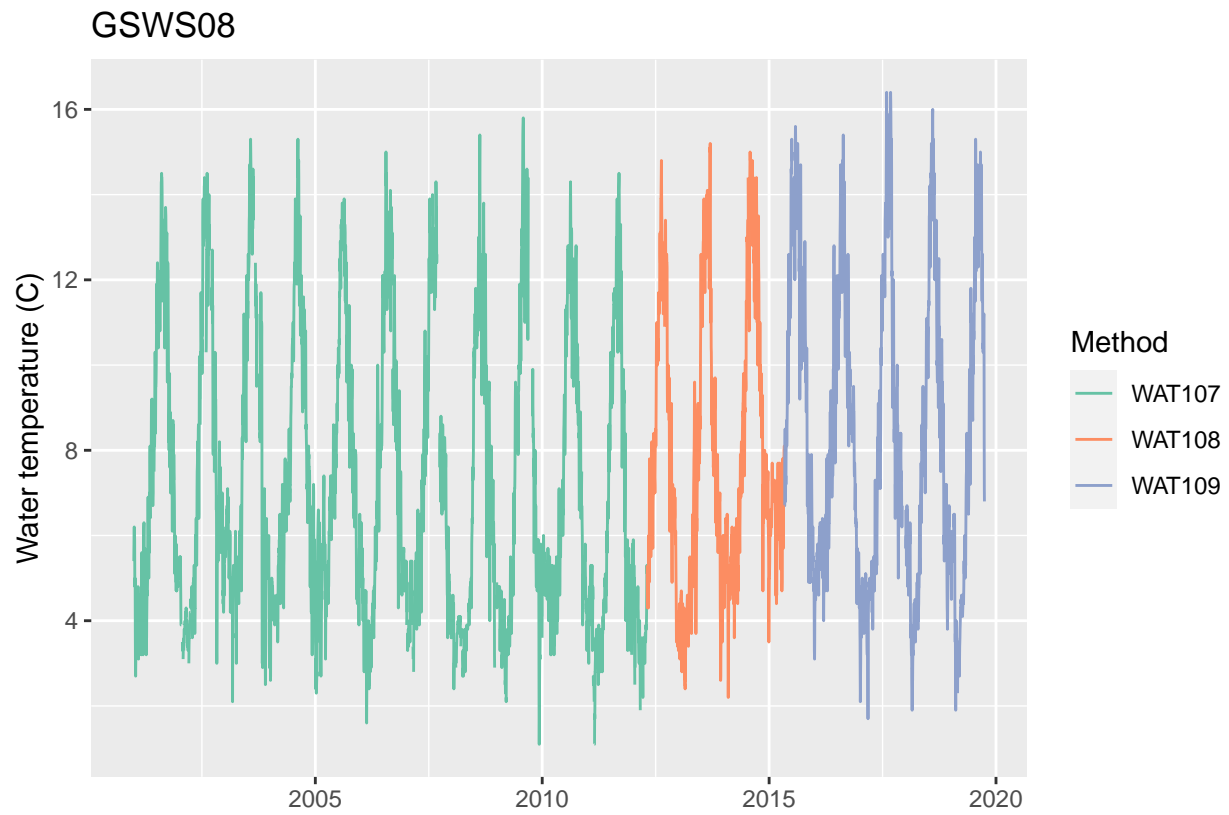


GSWS06

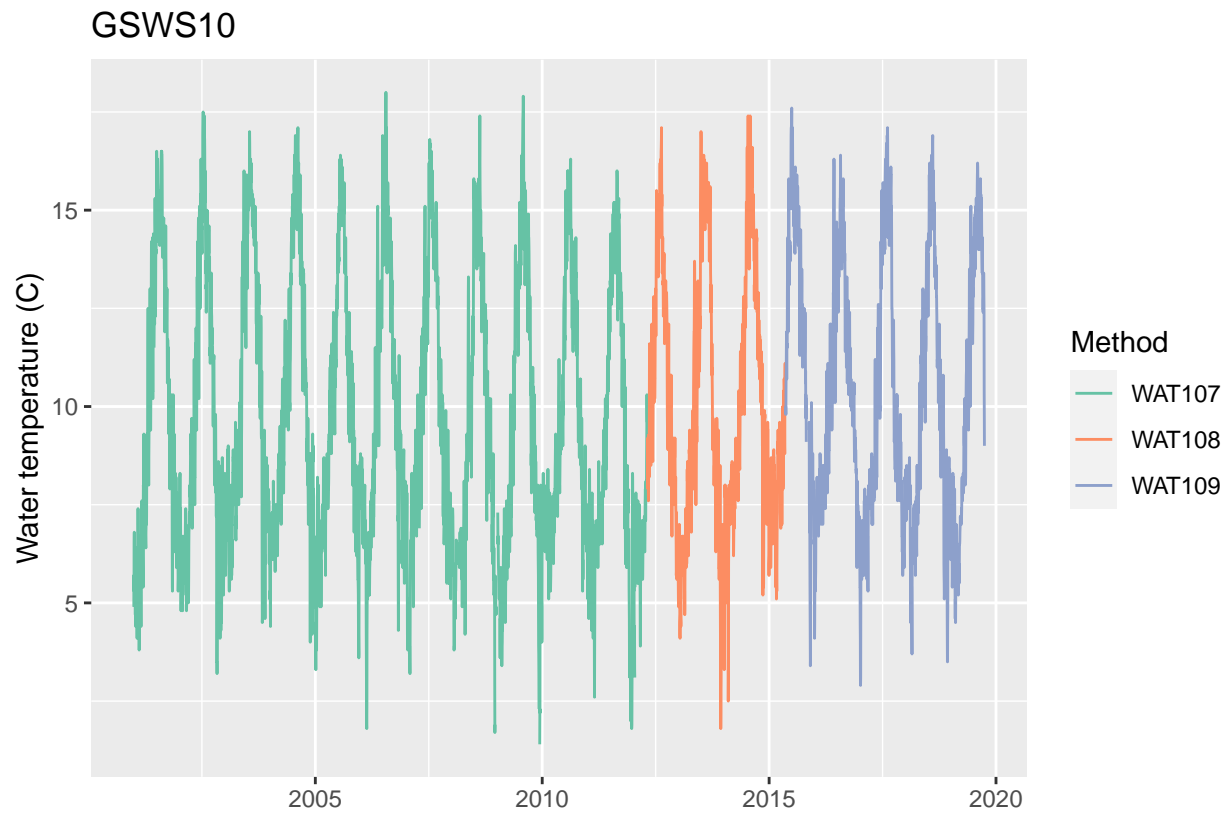


GSWS07



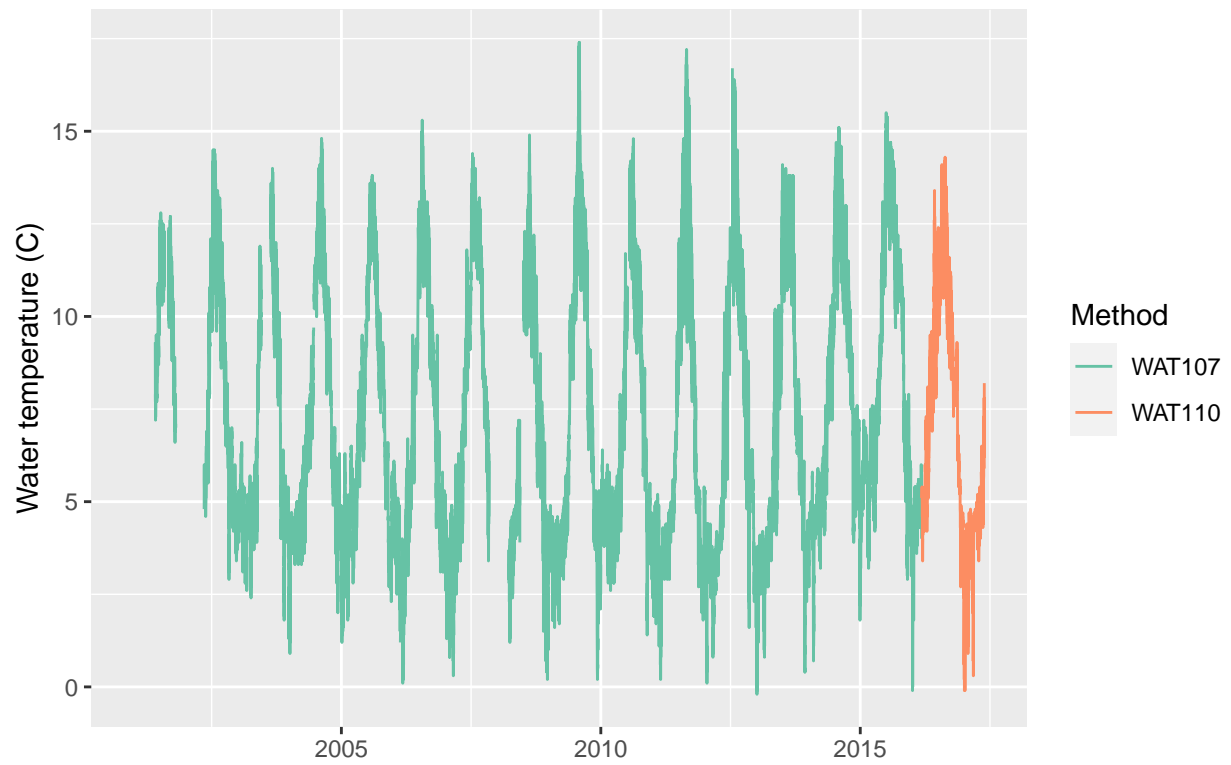


```
## Warning: Removed 4 row(s) containing missing values (geom_path).
```



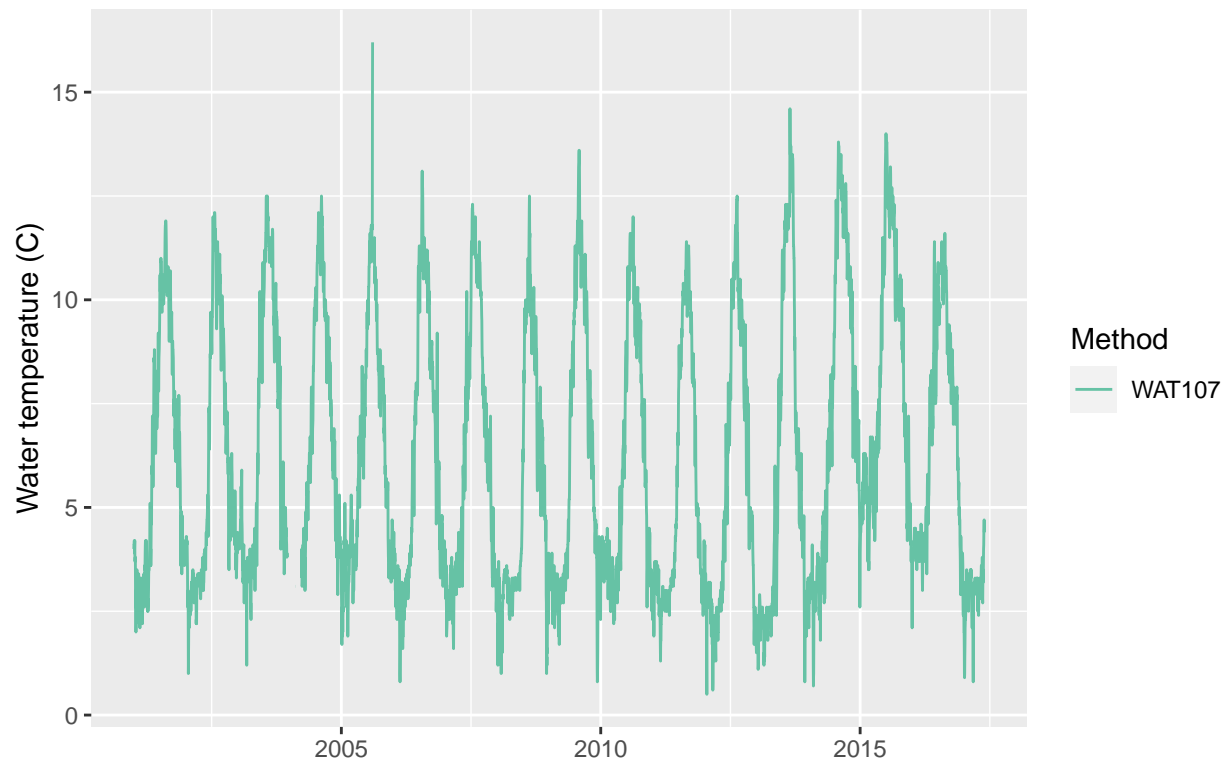
```
## Warning: Removed 4 row(s) containing missing values (geom_path).
```

## TSLOMA

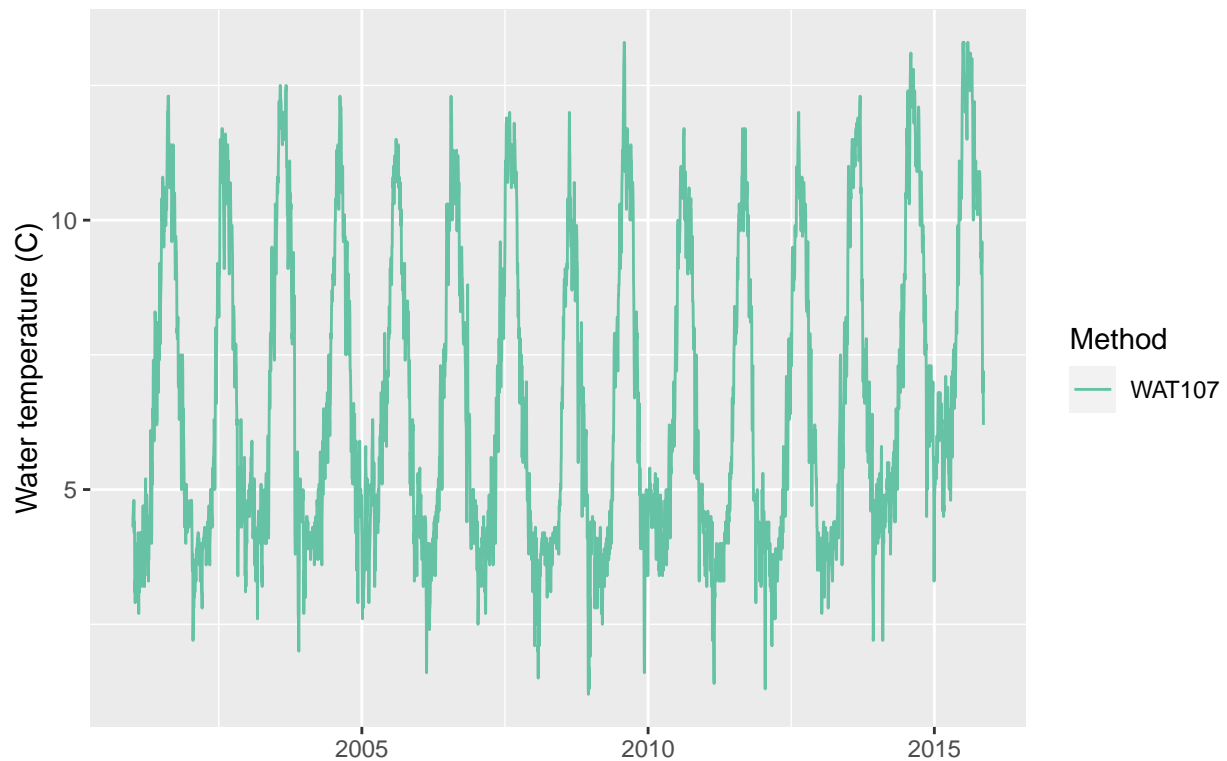


```
## Warning: Removed 1 row(s) containing missing values (geom_path).
```

## TSLOOK



## TSMCRA



dplyrs `group_walk()` function (this both creates and displays the plots):

```
streams %>%  
  group_by(sitecode) %>%  
  group_walk(~ print(plot_temperature(.x, .y)))
```

A for loop:

```
sitecodes <- unique(streams$sitecode)  
for (site in sitecodes){  
  site_data <- filter(streams,  
    sitecode == site)  
  site_data %>%  
    plot_temperature(site) %>%  
    print()  
}
```

Or alternatively save plots as PNG files. With purrrs `walk2()`

```
streams_with_plots %>%  
  mutate(  
    plot_files = paste0("plots/", sitecode, ".png")  
  ) %>%  
  with(walk2(plot_files, plots, ggsave))
```

```
## Saving 6.5 x 4.5 in image
```



```
## Warning: Removed 5 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 1 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 2 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 2 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 1 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image
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## Warning: Removed 4 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 4 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image

## Warning: Removed 1 row(s) containing missing values (geom_path).

## Saving 6.5 x 4.5 in image
```