

# Iterations and Functionals

Stat 220

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# Why repeat ourselves?

```
tinydata <- tribble(  
  ~case, ~x, ~y, ~z,  
  "a", 5, 3, -2,  
  "b", 7, 1, -5,  
  "c", 9, 12, -3  
)
```

```
tinydata  
# A tibble: 3 × 4  
  case      x      y      z  
  <chr> <dbl> <dbl> <dbl>  
1 a         5      3     -2  
2 b         7      1     -5  
3 c         9     12     -3
```

## Find the mean of each columns

```
mean(tinydata$x)  
[1] 7
```

```
mean(tinydata$y)  
[1] 5.333333
```

```
mean(tinydata$z)  
[1] -3.333333
```

# Iteration

Iteration is the process of repeating the same action over and over again

Multiple ways to do in R

- **loops** using `for`, `while`, etc
- **vectorized** functions that apply the same function to every element of a vector
- **functional** functions that apply the same function to elements in a vector, matrix, data frame, or list

# for loops

A way to iterate through a series of items stored as data object in R.

```
items <- c("grapes", "bananas", "chocolate", "bread")
for(i in items){
  print(i)
}
[1] "grapes"
[1] "bananas"
[1] "chocolate"
[1] "bread"
```

```
i <- items[1]
print(i)
[1] "grapes"
```

```
i <- items[2]
print(i)
[1] "bananas"
```

# for loop components

the for() function which we use to specify

- what object we're drawing from and
- what object we are writing to.

```
for( i   in  items  )  
    ^       ^  
    |       |___ object we are drawing from  
    |  
obj. we write each item to
```

# for loop components

## The brackets {}

- Inside the brackets we house the code that is going to happen each iteration.

```
for( i in items ){  
    | ~~~~~|  
    | ~~~~~|  
    | ~~~~~| code we need perform on each iteration.  
    | ~~~~~|  
    | ~~~~~|  
}
```

# for() loops and storing output

```
letters
[1] "a" "b" "c" "d" "e" "f" "g" "h" "i" "j" "k" "l" "m" "n" "o" "p" "q" "r" "s"
[20] "t" "u" "v" "w" "x" "y" "z"
```

```
basket <- rep(NA,10) # numeric vector of length 3
basket
[1] NA NA NA NA NA NA NA NA NA NA
```

```
# Each loop, we store the output of some code.
for(i in 1:10){
  basket[i] <- str_glue(letters[i],letters[i+1])
}
basket
[1] "ab" "bc" "cd" "de" "ef" "fg" "gh" "hi" "ij" "jk"
```

# For loops tidydata

- Let's iterate calculation of column means:

```
my_means <- rep(NA, 3)
my_means
[1] NA NA NA
```

```
for (i in 1:3) { # three columns to get the mean for
  my_means[i] <- mean(tinydata[[i+1]]) # mean of col. i+1 (skip col. 1)
}
my_means
[1] 7.000000 5.333333 -3.333333
```



# For loops: preallocation output space

- About 12 seconds without preallocation and less than a second with (elapsed time).

## Without preallocation

```
system.time({  
  output <- NULL  
  for (i in 1:1000000) {  
    output <- c(output, i)  
  }  
})  
   user  system elapsed  
11.924   0.016  11.947
```

## With preallocation

```
system.time({  
  output <- rep(NA, 1000000)  
  for (i in 1:1000000) {  
    output[i] <- i  
  }  
})  
   user  system elapsed  
0.008   0.000   0.008
```

# For loops: index vector

- `seq_along(df)` index based on columns of data frame

```
seq_along(tinydata)
[1] 1 2 3 4
```

- Another common way of indexing

```
1:nrow(tinydata)
[1] 1 2 3
```

- Along the columns

```
1:ncol(tinydata)
[1] 1 2 3 4
```

# For loop with ifelse

```
my_means <- rep(NA, ncol(tinydata))

for (i in seq_along(tinydata)){ # iterate over all columns
  my_means[i] <- ifelse(is.numeric(tinydata[[i]]), mean(tinydata[[i]]), NA)
}
```

```
my_means
[1]      NA  7.000000  5.333333 -3.333333
```

# Function for conditional evaluation

- if x is numeric then standardize, else just return x

```
standardize <- function(x, ...){      # ... used for arbitrary number of arguments
  if (is.numeric(x)){                 # condition
    (x - mean(x, ...))/sd(x, ...)      # if TRUE, standardize
  } else{                             # else (FALSE)
    x                                  # return x unchanged
  }
}
```

```
standardize(c(1,2,3,4))
[1] -1.1618950 -0.3872983  0.3872983  1.1618950
```

```
standardize(c("a", "b", "2", NA), na.rm = TRUE)
[1] "a" "b" "2" NA
```

# Standardizing tinydata

- Allocate storage in a new data frame:

```
scaled_tinydata <- tinydata %>%  
  mutate(  
    x = NA,  
    y = NA,  
    z = NA  
  )
```

```
scaled_tinydata  
# A tibble: 3 × 4  
  case  x      y      z  
  <chr> <lgl> <lgl> <lgl>  
1 a     NA     NA     NA  
2 b     NA     NA     NA  
3 c     NA     NA     NA
```

# Standardizing tinydata

- For loop for iteration:

```
for (i in seq_along(tinydata)){  
  scaled_tinydata[, i] <- standardize(tinydata[[i]])  
}
```

```
scaled_tinydata  
# A tibble: 3 × 4  
  case      x      y      z  
  <chr> <dbl> <dbl> <dbl>  
1 a      -1 -0.398  0.873  
2 b       0 -0.740 -1.09  
3 c       1  1.14   0.218
```

# Vectorization

- A vectorized function will apply the same operation (function) to each element of a vector.
  - avoid loops by applying operations to each element of a vector

# Vectorization

```
x <- c(10,20,30,40)
log10(x)      # log10 is a vectorized function
[1] 1.000000 1.301030 1.477121 1.602060
```

- The for loop version

```
out <- rep(NA, 4)
for (i in 1:4)
  { out[i] <- log10(x[i]) }
out
[1] 1.000000 1.301030 1.477121 1.602060
```



## Your Turn 1

Load the `palmerpenguins` package. Write a `for` loop that calculates the mean of the numeric variables in the `penguins` data set and stores the means in a named vector.

```
Rows: 344
Columns: 8
$ species      <fct> Adelie, Adelie, Adelie, Adelie, Adelie, Adelie, Adel...
$ island       <fct> Torgersen, Torgersen, Torgersen, Torgersen, Torgersen, Torgerse...
$ bill_length_mm <dbl> 39.1, 39.5, 40.3, NA, 36.7, 39.3, 38.9, 39.2, 34.1, ...
$ bill_depth_mm <dbl> 18.7, 17.4, 18.0, NA, 19.3, 20.6, 17.8, 19.6, 18.1, ...
$ flipper_length_mm <int> 181, 186, 195, NA, 193, 190, 181, 195, 193, 190, 186...
$ body_mass_g   <int> 3750, 3800, 3250, NA, 3450, 3650, 3625, 4675, 3475, ...
$ sex          <fct> male, female, female, NA, female, male, female, male...
$ year         <int> 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007, 2007...
```

04:00

# Functionals

- A **functional** function will apply the same operation (function) to each element of a vector, matrix, data frame or list.
- base-R: `apply` family of commands
  - `purrr` package: `map` family of commands



# apply family of commands

- R has a family of commands that apply a function to different parts of a vector, matrix or data frame.

## **lapply(X, FUN):**

applies the *FUN* to each element in the vector/list *X*. Returns a list with length equal to that of the vector

## **sapply:**

works like `lapply` but returns a vector (so *FUN* can only return one value)

# apply family of commands

- R has a family of commands that apply a function to different parts of a vector, matrix or data frame.

**apply(matrix, MARGIN, FUN):**

applies the function *FUN* to the matrix. *MARGIN* given (1=row, 2=column, c(1,2)=rows and cols). Returns an atomic **vector** or **matrix**

**tapply(x,y,FUN):**

applies *FUN* to atomic vector (variable) *x* for each group in categorical variable *y*. Returns an atomic **vector** with a `dim names` attribute

# purrr package

powerful package for iteration with the same functionality as apply commands, but more readable (according to Hadley)



- `map(.x, .f)` maps the function `.f` to elements in the vector/list `.x`

# lapply tiny example

```
lapply(tinydata, FUN = mean)
$case
[1] NA

$x
[1] 7

$y
[1] 5.333333

$z
[1] -3.333333
```

- a 3x4 data frame is **summarized** in a list of length 4.

- R sees tinydata as a list whose elements are column vectors (variables)
- the FUN is applied to each list element
- a list is returned
- length is the number of variables in the data frame

# map

In purrr, the map function is equivalent to lapply

```
library(purrr)
map(tinydata, .f = mean)
$case
[1] NA

$x
[1] 7

$y
[1] 5.333333

$z
[1] -3.333333
```

# sapply tiny example

Output is an atomic vector (simplify)

```
sapply(tinydata, FUN = mean)
      case      x      y      z
      NA  7.000000  5.333333 -3.333333
```

- a 3x4 data frame is **summarized** in a vector of length 4.



# map\_dbl

map\_dbl is equivalent to sapply

```
map_dbl(tinydata, .f = mean)
```

case	x	y	z
NA	7.000000	5.333333	-3.333333

# map\_df

map\_df returns a data frame instead of a vector

```
map_df(tinydata, .f = mean)
# A tibble: 1 × 4
  case      x      y      z
<dbl> <dbl> <dbl> <dbl>
1     NA      7  5.33 -3.33
```

- No equivalency in base-R apply!

# Iterate or dplyr?!

- `summarize_all`, `summarize_if`, `summarize_at` are all options that apply `.fun`s to **columns** of a data frame
- `if` option needs a logical function that determines which **columns** to apply the `.fun`s to

```
tinydata %>%  
  summarize_if(is.numeric, .fun = mean)  
# A tibble: 1 × 3  
      x      y      z  
  <dbl> <dbl> <dbl>  
1      7  5.33 -3.33
```

# functionals: single function that mutates

standardize function gives us a list of standardized values

```
lapply(tinydata, FUN = standardize)
$case
[1] "a" "b" "c"

$x
[1] -1  0  1

$y
[1] -0.3982161 -0.7395442  1.1377602

$z
[1]  0.8728716 -1.0910895  0.2182179
```

- a 3x4 data frame is **mutated** to a list of 4 vectors of length 3 each

# lapply tiny example

Using `dplyr::bind_cols` converts the list to a data frame with variables equal to list entries

```
lapply(tinydata, FUN = standardize) %>%
```

```
  bind_cols()
```

```
# A tibble: 3 × 4
```

	case	x	y	z
	<chr>	<dbl>	<dbl>	<dbl>
1	a	-1	-0.398	0.873
2	b	0	-0.740	-1.09
3	c	1	1.14	0.218

# map\_df

In purrr, the map\_df is equal to lapply + bind\_cols:

```
map_df(tinydata, .f = standardize)
```

```
# A tibble: 3 × 4
```

	case	x	y	z
	<chr>	<dbl>	<dbl>	<dbl>
1	a	-1	-0.398	0.873
2	b	0	-0.740	-1.09
3	c	1	1.14	0.218

- a 3x4 data frame is mutated to **standardized** 3x4 data frame

# Iterate or dplyr?!

- `mutate_all`, `mutate_if`, `mutate_at` are all options that apply `.fun`s to columns of a data frame
- if option needs a logical function that determines which columns to apply the `.fun`s to

```
tinydata %>%  
  mutate_if(is.numeric, .fun = standardize)  
# A tibble: 3 × 4  
  case      x      y      z  
  <chr> <dbl> <dbl> <dbl>  
1 a      -1 -0.398  0.873  
2 b       0 -0.740 -1.09  
3 c       1  1.14   0.218
```

# applying multiple functions

- Let's get the 0.1 and 0.9 quantile for variables in `tinydata`

```
quantile(tinydata$x, probs = c(.1, .9))  
10% 90%  
5.4 8.6
```

```
quantile(tinydata$y, probs = c(.1, .9))  
10% 90%  
1.4 10.2
```

```
quantile(tinydata$z, probs = c(.1, .9))  
10% 90%  
-4.6 -2.2
```

- the function output is a vector of length 2 (same lengths as `probs`)



# map\_df: getting quantiles

```
tinydata %>%  
  select_if(is.numeric) %>%    # only numeric columns  
  map_df(  
    .f = quantile,    # function to apply to cols  
    probs = c(.1, .9)) # extra function arguments  
# A tibble: 3 × 2  
  `10%` `90%`  
  <dbl> <dbl>  
1    5.4    8.6  
2    1.4   10.2  
3   -4.6   -2.2
```

# map\_df: getting quantiles

Can use `.id` to record the variable names from `tinydata`:

```
tinydata %>%  
  select_if(is.numeric) %>%  
  map_df(  
    .f = quantile,  
    probs = c(.1, .9),  
    .id = "variable")  
# A tibble: 3 × 3  
  variable `10%` `90%`  
  <chr>    <dbl> <dbl>  
1 x        5.4   8.6  
2 y        1.4  10.2  
3 z       -4.6  -2.2
```

## map\_df options

There are two types of map\_df

- map\_dfr: which row binds the list created by map
  - entries in the list are rows in the data frame
- map\_df: which column binds the list created by map
  - entries in the list are columns in the data frame

# Iterate or dplyr?!

- `summarize_all`, `summarize_if`, `summarize_at` can work with functions like `quantile` that return multiple values.
- the **form** of the output is a transposed version of `map_df`

```
tinydata %>%  
  summarize_if(is.numeric, .funs = quantile, probs = c(.1, .9))  
# A tibble: 2 × 3  
      x      y      z  
  <dbl> <dbl> <dbl>  
1   5.4   1.4  -4.6  
2   8.6  10.2  -2.2
```

- **rows** = 0.1 and 0.9 quantiles
- **cols** = variables

# Iterate or dplyr?!

- We need to manually add a percentile variable to help us ID the value in each row

```
tinydata %>%  
  summarize_if(is.numeric, .funs = quantile, probs = c(.1, .9)) %>%  
  mutate(percentile = c(10,90))  
# A tibble: 2 × 4  
      x      y      z percentile  
  <dbl> <dbl> <dbl>      <dbl>  
1   5.4   1.4  -4.6         10  
2   8.6  10.2  -2.2         90
```

# Example: Energy Data

- Recall the wide version of the energy data:

```
energy %>% select(dayWeek, `Center_for_Mathematics_&_Computing`) %>% glimpse
Rows: 35,129
Columns: 2
$ dayWeek                <fct> Tues, Tues, Tues, Tues, Tues, Tue...
$ `Center_for_Mathematics_&_Computing` <dbl> 14.3750, 14.0625, 14.3750, 14.062...
```

# Quantiles from wide data

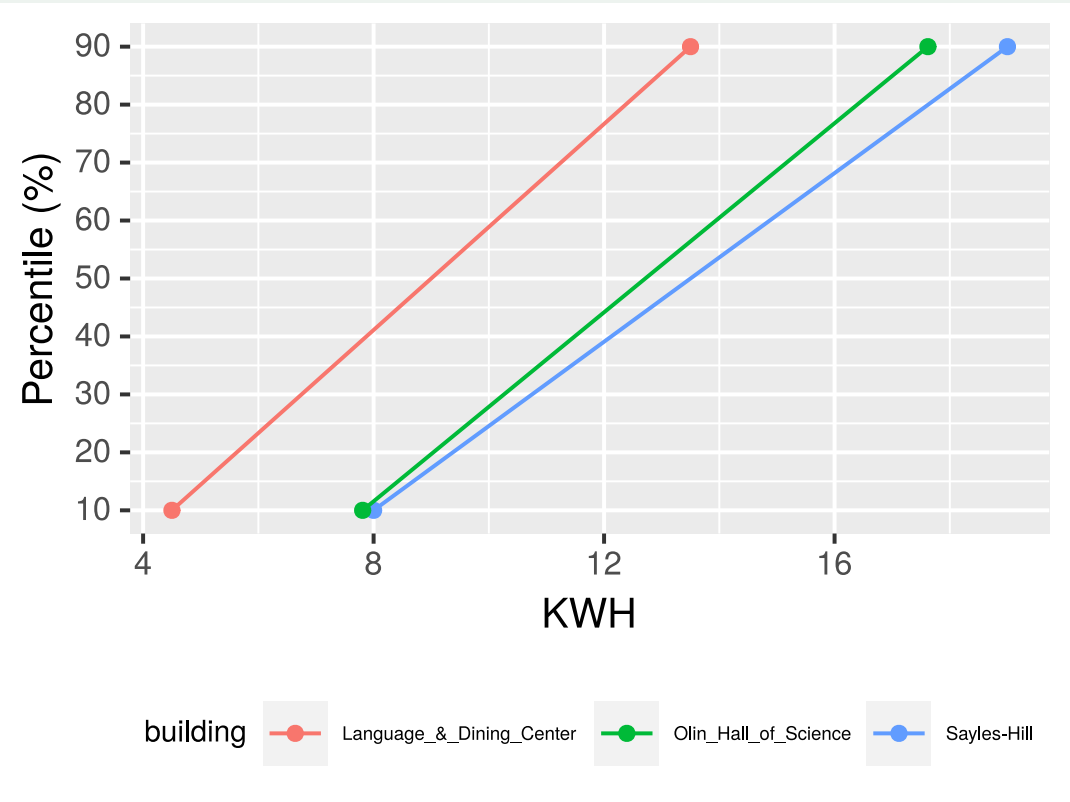
- Let's get 0.1 and 0.9 quantiles for 3 buildings:

```
energy_quant <- energy %>%  
  select("Sayles-Hill" , "Language_&_Dining_Center", "Olin_Hall_of_Science") %>%  
  map_df(  
    .f = quantile,  
    probs = c(.1, .9),  
    na.rm = TRUE,  
    .id = "building")
```

```
# A tibble: 3 × 3  
  building      `10%` `90%`  
  <chr>      <dbl> <dbl>  
1 Sayles-Hill      8     19  
2 Language_&_Dining_Center  4.5   13.5  
3 Olin_Hall_of_Science    7.81  17.6
```

# Plot of Quantiles

```
energy_quant %>%  
  mutate(percentile = parse_number(percentile))  
  ggplot(aes(y = percentile, x = value)) +  
  geom_point() +  
  geom_line(aes(group=building)) +  
  labs(y="Percentile (%)", x="KWH") +  
  scale_y_continuous(breaks=seq(10, 90, 10)) +  
  theme(legend.position="bottom",  
        legend.text = element_text(size=12),  
        legend.title=element_text(size=14))
```





# Quantiles from wide data

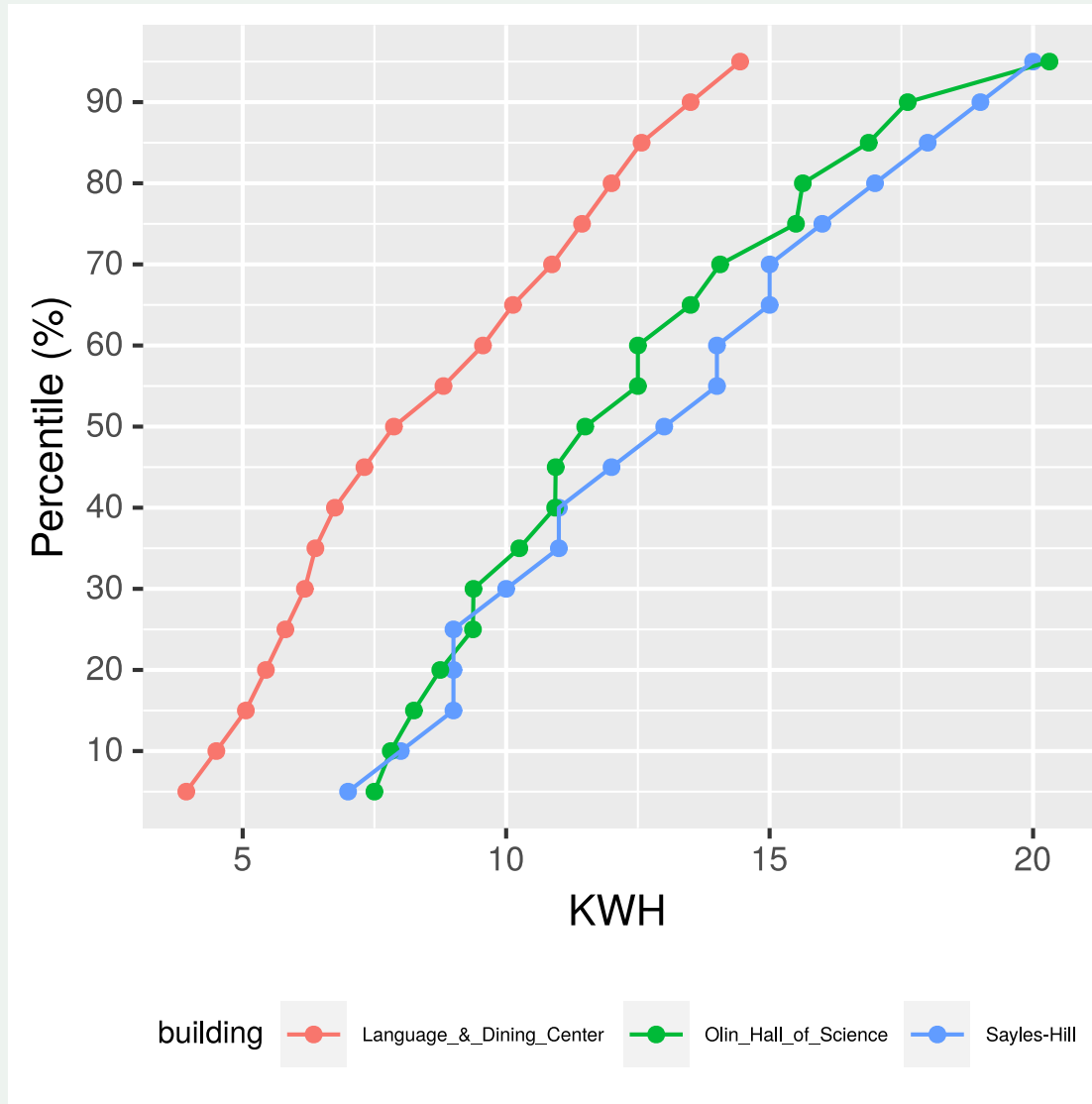
- create a vector of quantile probs:

```
p <- seq(0.05,0.95, by = .05) # every 5th quantile
energy_quant <- energy %>%
  select("Sayles-Hill" ,"Language_&_Dining_Center", "Olin_Hall_of_Science") %>%
  map_df(
    .f = quantile,
    probs = p,
    na.rm = TRUE,
    .id = "building") %>%
  pivot_longer(
    names_to = "percentile",
    values_to = "value",
    cols = 1 + 1:length(p), # quantiles start in col 2
  )
```

# Quantiles from wide data

```
energy_quant
# A tibble: 57 × 3
  building    percentile value
  <chr>        <chr>     <dbl>
1 Sayles-Hill 5%         7
2 Sayles-Hill 10%        8
3 Sayles-Hill 15%        9
4 Sayles-Hill 20%        9
5 Sayles-Hill 25%        9
6 Sayles-Hill 30%       10
7 Sayles-Hill 35%       11
8 Sayles-Hill 40%       11
9 Sayles-Hill 45%       12
10 Sayles-Hill 50%       13
# ... with 47 more rows
```

# Quantiles from wide data



# Quantiles from long data

- What if we have the long version of this data?

```
energy_long <- energy %>%  
  pivot_longer(  
    names_to = "building",  
    values_to = "energyKWH",  
    cols = 9:90) %>%  
  filter(building %in% c("Sayles-Hill" ,  
                        "Language_&_Dining_Center",  
                        "Olin_Hall_of_Science") )
```

# Quantiles from long data

**Goal:** get quantiles for every `building` and `dayWeek`

- We don't have many columns of measurements to apply a function to
- We have groups (`building` and `month`) that we need to summarize with quantile (more than one output value)

# Quantiles from long data

- Let's get quantiles for every building and day of the week:

```
energy_long_quant <- energy_long %>%  
  group_by(building, dayWeek) %>%  
  summarize(value = quantile(energyKWH, probs = c(.1, .9), na.rm = TRUE))
```

```
# A tibble: 42 × 3  
# Groups:   building, dayWeek [21]  
  building          dayWeek value  
  <chr>          <fct>    <dbl>  
1 Language_&_Dining_Center Mon      4.88  
2 Language_&_Dining_Center Mon     14.1  
3 Language_&_Dining_Center Tues      4.5  
4 Language & Dining Center Tues     13.9
```

# Quantiles from long data

- Need to add a percentile

```
energy_long_quant <- energy_long_quant %>%  
  mutate(percentile = c(10,90))
```

```
energy_long_quant  
# A tibble: 42 × 4  
# Groups:   building, dayWeek [21]  
  building          dayWeek value percentile  
  <chr>          <fct>    <dbl>      <dbl>  
1 Language_&_Dining_Center Mon      4.88        10  
2 Language_&_Dining_Center Mon     14.1        90  
3 Language & Dining Center  Tues      4.5        10
```

# Quantiles from long data

For all quantiles in `p <- seq(0.05,0.95, by = 0.05))`

```
energy_long_quant <- energy_long %>%  
  group_by(building, dayWeek) %>%  
  summarize(value = quantile(energyKWH, probs = p, na.rm = TRUE)) %>%  
  mutate(percentile = 100*p)
```

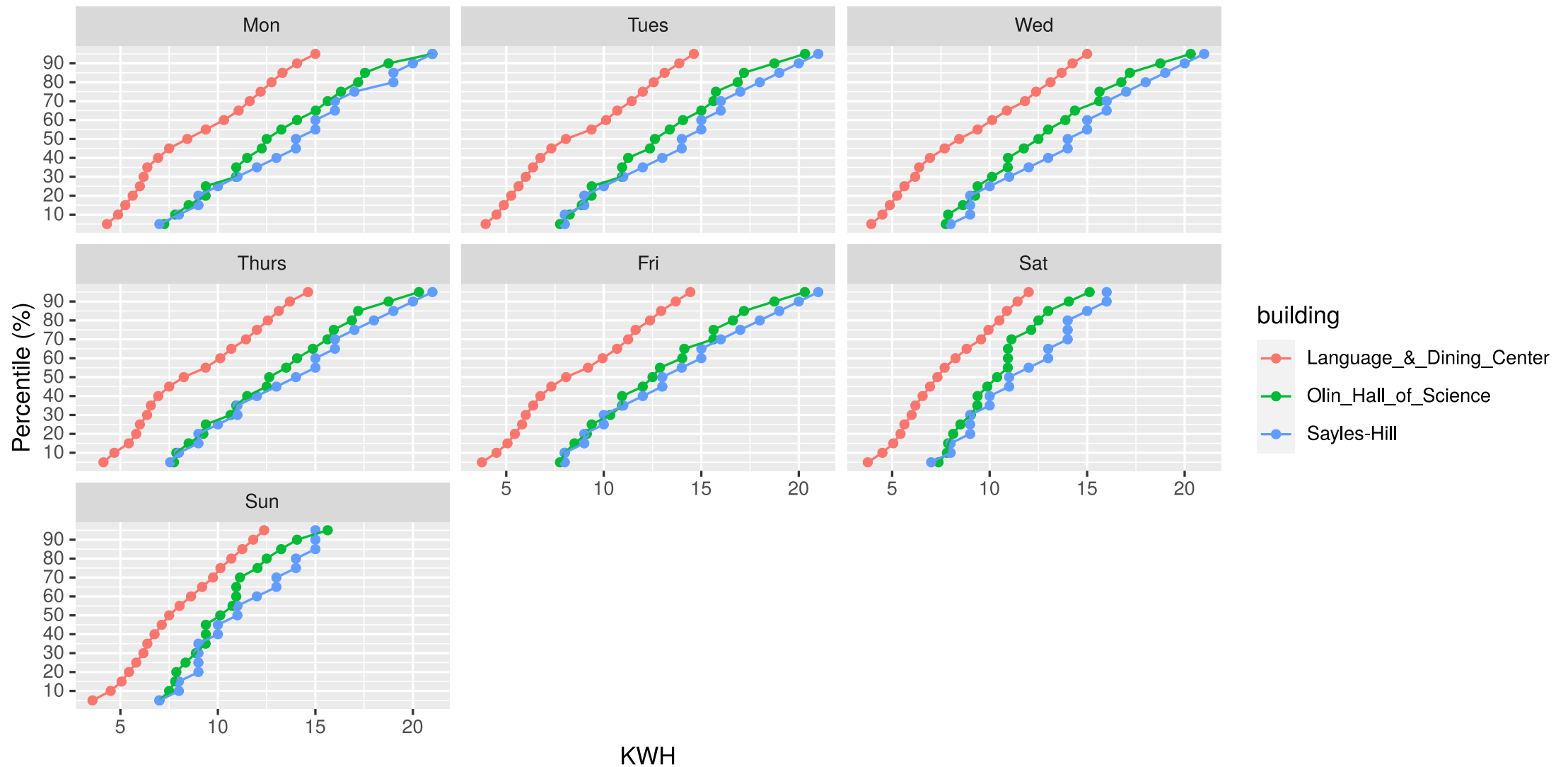
```
# A tibble: 399 × 4  
# Groups:   building, dayWeek [21]  
  building      dayWeek value percentile  
  <chr>      <fct>    <dbl>      <dbl>  
1 Language_&_Dining_Center Mon      4.31         5  
2 Language_&_Dining_Center Mon      4.88        10  
3 Language_&_Dining_Center Mon      5.25        15  
4 Language & Dining Center Mon      5.63        20
```



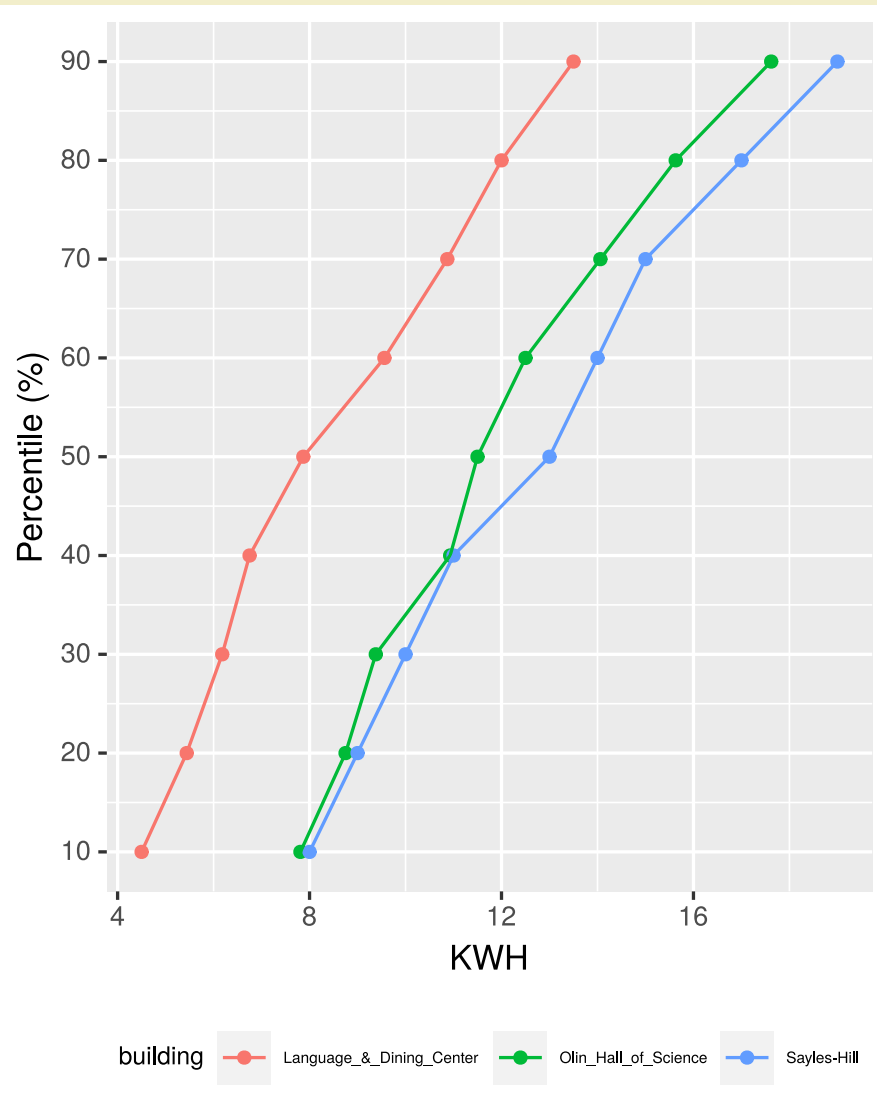
## Quantiles from long data

```
energy_long_quant %>%  
  ggplot(aes(y = percentile, x = value, color=building)) +  
  geom_point() +  
  geom_line(aes(group=building)) +  
  labs(y="Percentile (%)", x="KWH") +  
  scale_y_continuous(breaks=seq(10,90,by=10)) +  
  facet_wrap(~dayWeek)
```

# Quantiles from long data



## Your Turn 2



Follow the prompts to plot the quantiles of energy consumption for the buildings Sayles-Hill, Language\_&\_Dining\_Center, Olin\_Hall\_of\_Science.

05:00