

# Data Visualization

## Part II

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# 1 Agenda

We are going to **visually** analyze two datasets and see if we can tell stories from the visuals.



```
Registered S3 method overwritten by 'printr':  
  method      from  
  knit_print.data.frame rmarkdown
```

## 2 Setting up

Let's first load the `ggplot2` package:

```
1 if (!require(ggplot2)) {  
2   install.packages("ggplot2") # install if not already installed  
3 }
```

Loading required package: `ggplot2`

```
1 library (ggplot2)
```

### 3 Titanic Survival



#### 3.1 Load the dataset

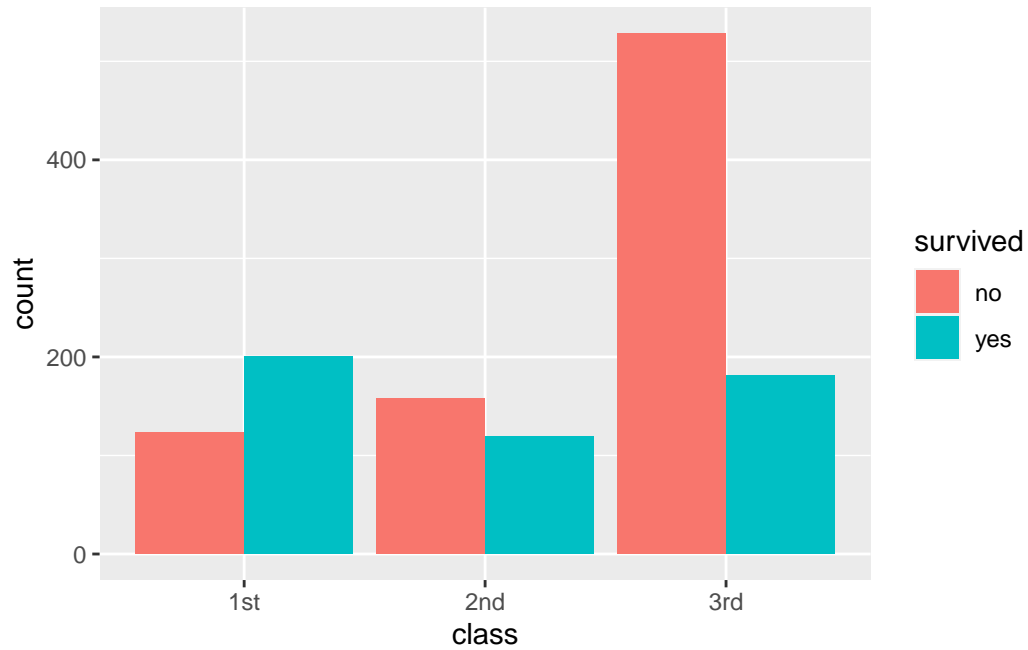
```
1 titanic = read.csv
  ↪ ("https://raw.githubusercontent.com/ahmedmoustafa/datasets/main/titanic/titanic.csv")
2 head(titanic)
```

name	survived	sex	age	class
Allen, Miss. Elisabeth Walton	yes	female	29.0000	1st
Allison, Master. Hudson Trevor	yes	male	0.9167	1st
Allison, Miss. Helen Loraine	no	female	2.0000	1st
Allison, Mr. Hudson Joshua Crei	no	male	30.0000	1st
Allison, Mrs. Hudson J C (Bessi	no	female	25.0000	1st
Anderson, Mr. Harry	yes	male	48.0000	1st

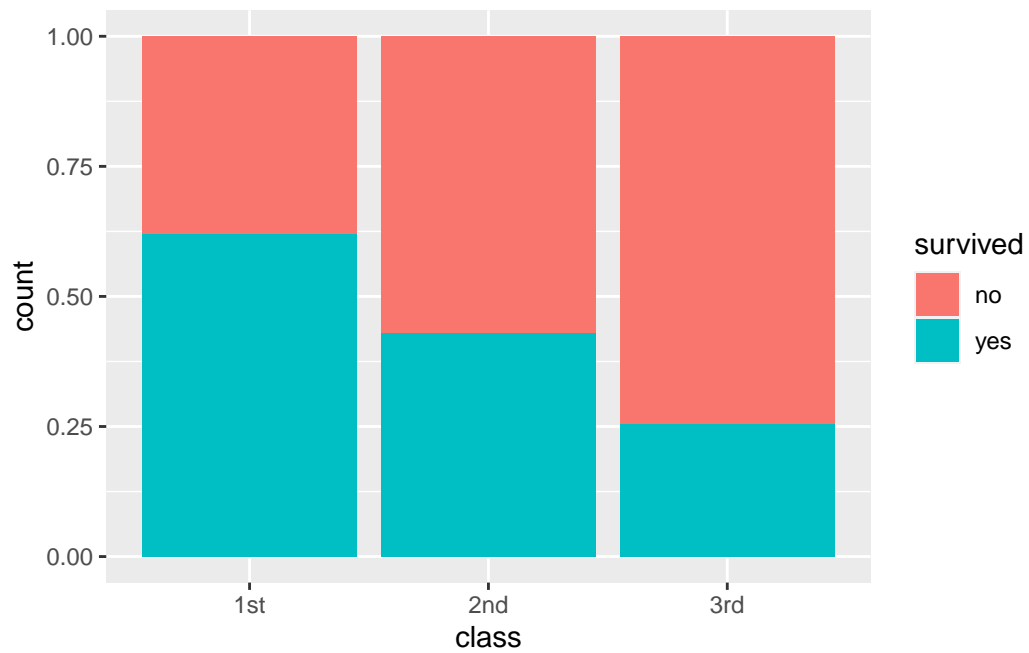
name	survived	sex	age	class
------	----------	-----	-----	-------

### 3.2 Survival by class

```
1 ggplot(titanic) +
2   geom_bar(aes(x = class, fill = survived), position = "dodge")
```

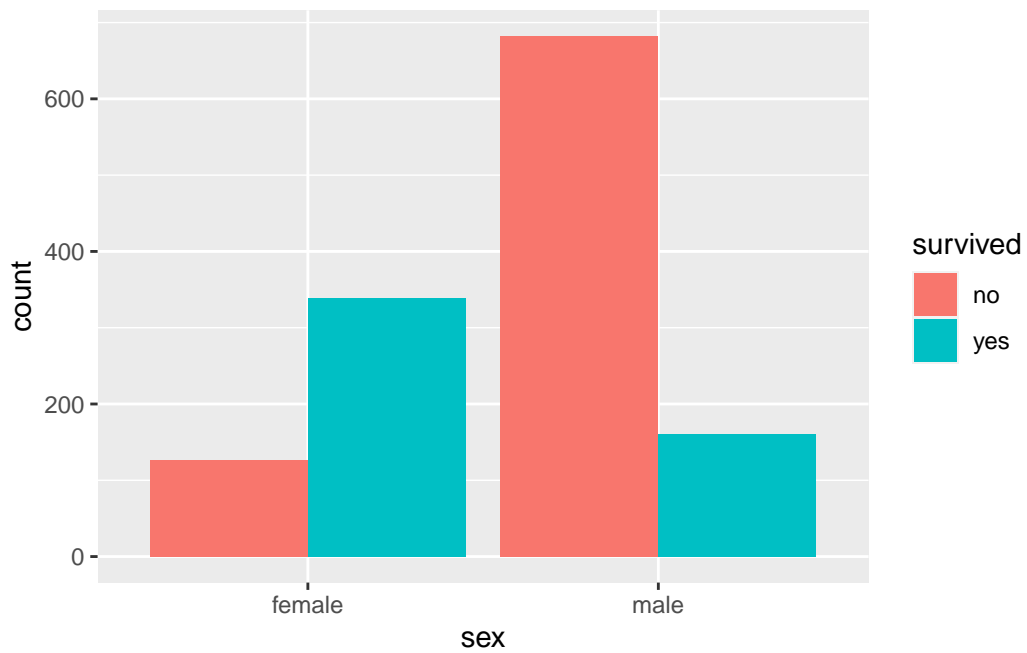


```
1 ggplot(titanic) +
2   geom_bar(aes(x = class, fill = survived), position = "fill")
```

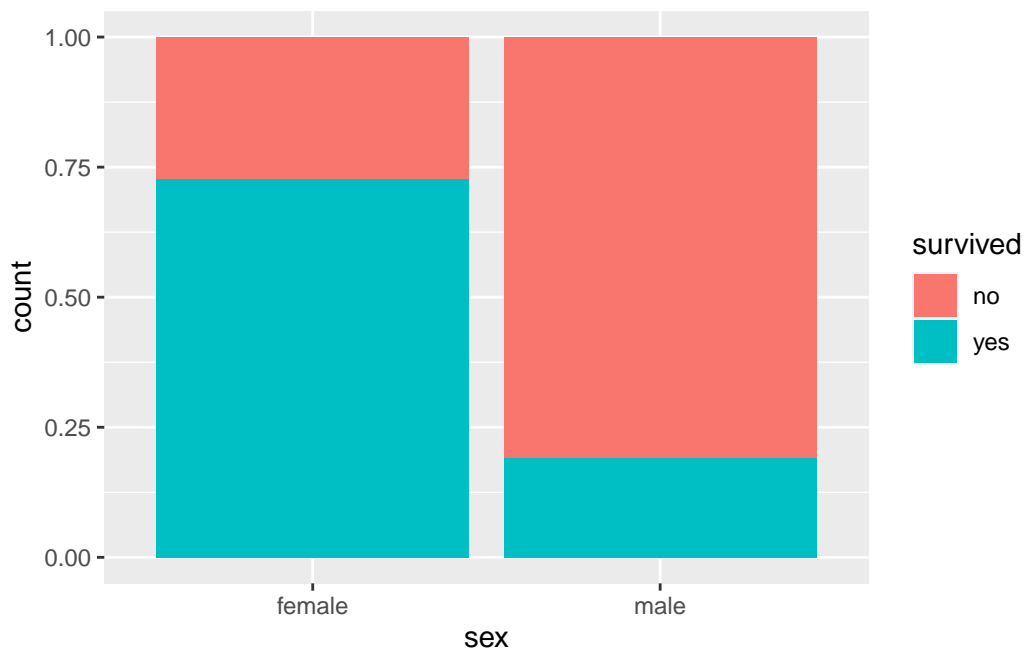


### 3.3 Survival by sex

```
1 ggplot(titanic) +  
2   geom_bar(aes(x = sex, fill = survived), position = "dodge")
```



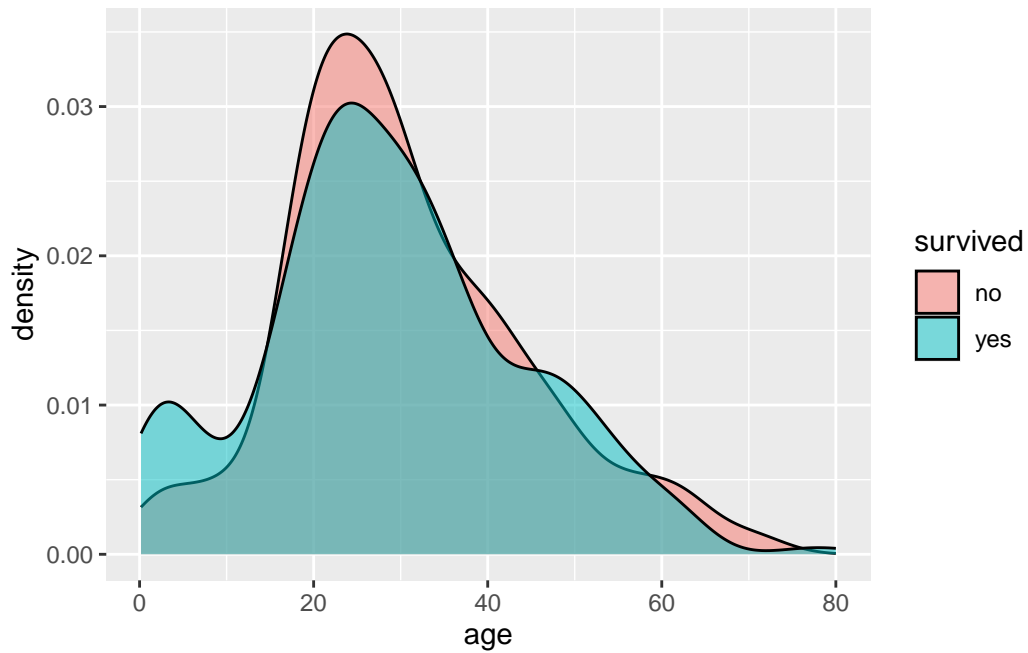
```
1 ggplot(titanic) +  
2   geom_bar(aes(x = sex, fill = survived), position = "fill")
```



### 3.4 Survival by age

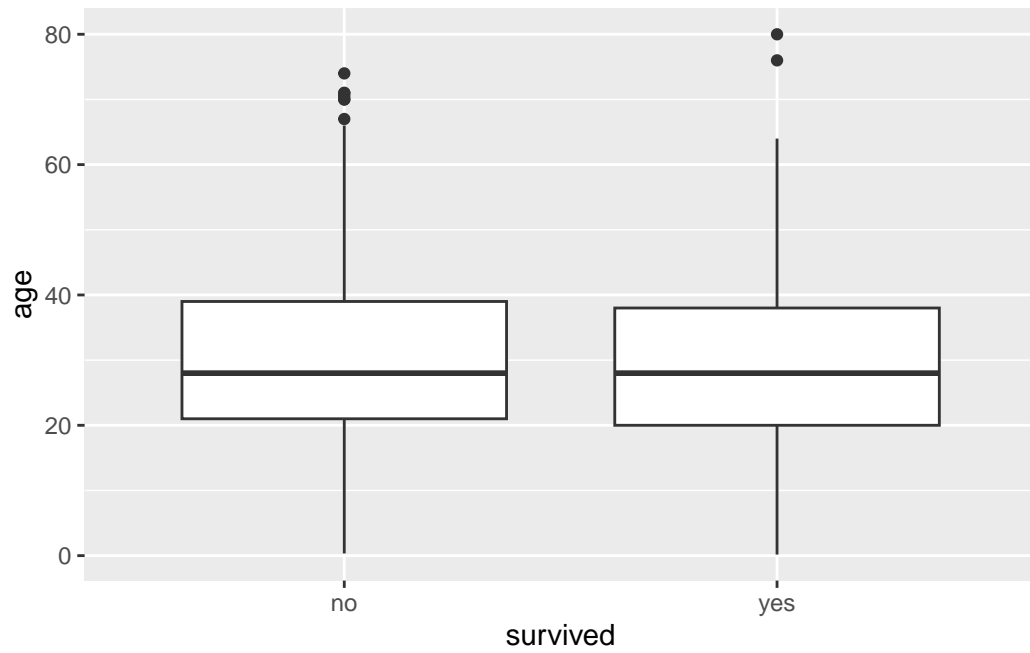
```
1 ggplot(titanic) +  
2   geom_density(aes(x = age, fill = survived), alpha = 0.5)
```

Warning: Removed 263 rows containing non-finite values (`stat\_density()`).



```
1 ggplot(titanic) +  
2   geom_boxplot(aes(x = survived, y = age))
```

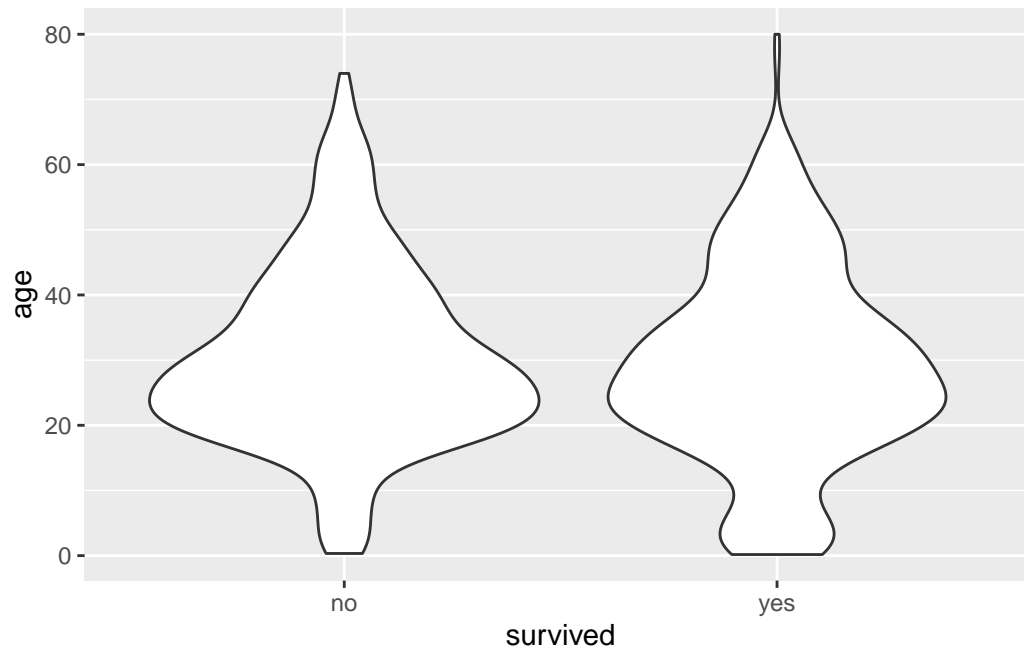
Warning: Removed 263 rows containing non-finite values (`stat\_boxplot()`).



```
1 ggplot(titanic) +  
2   geom_violin(aes(x = survived, y = age))
```

Warning: Removed 263 rows containing non-finite values (`stat\_ydensity()`).

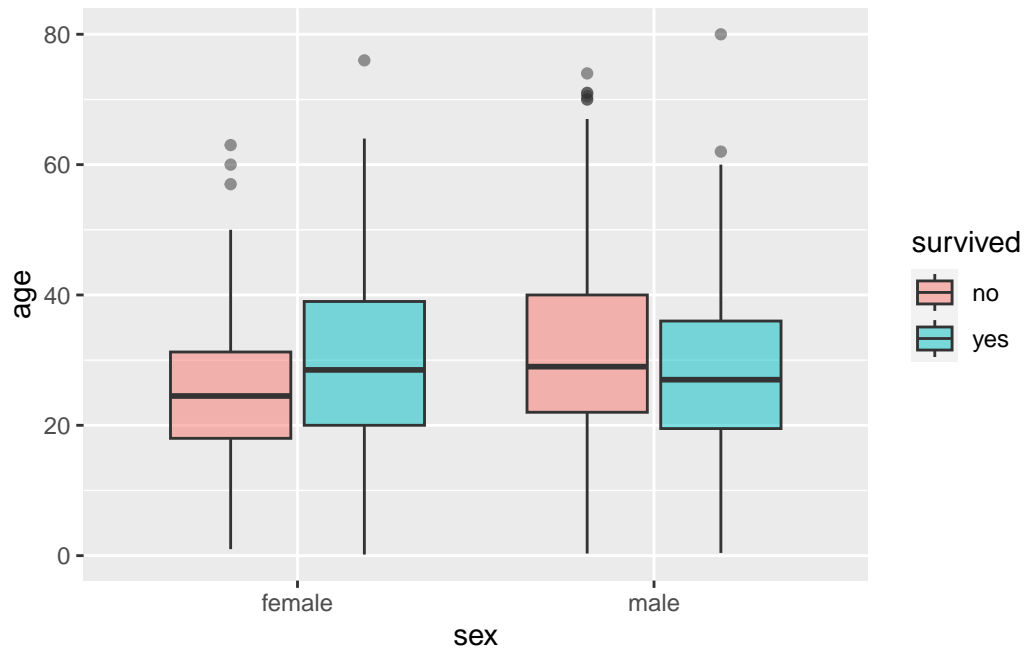




### 3.5 Survival by age & sex

```
1 ggplot(titanic) +  
2   geom_boxplot(aes(x = sex, y = age, fill = survived), alpha = 0.5)
```

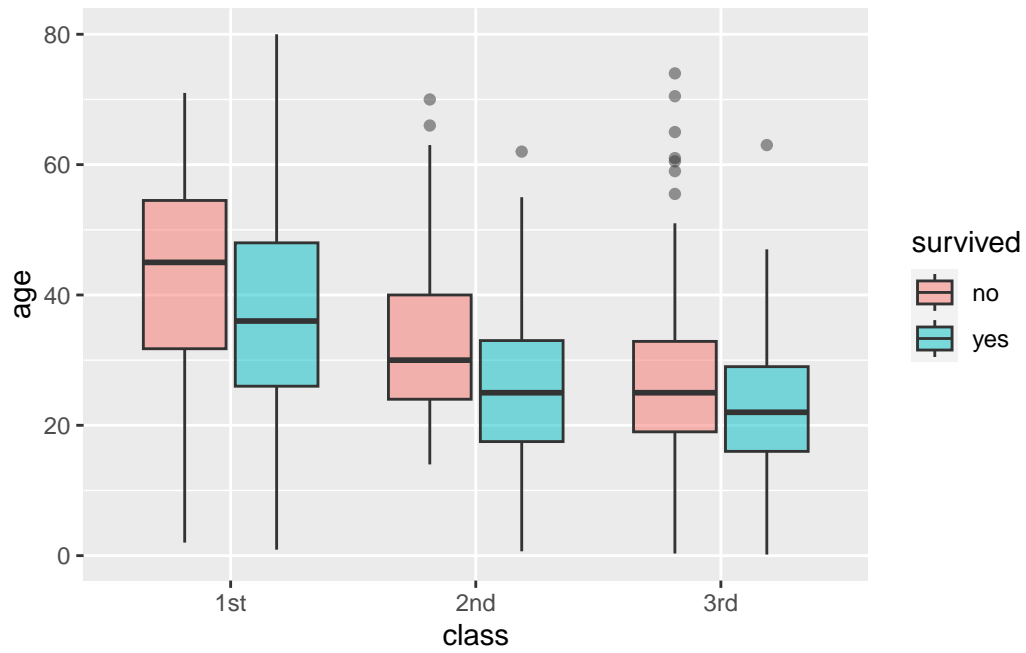
Warning: Removed 263 rows containing non-finite values (`stat\_boxplot()`).



### 3.6 Survival by class & age

```
1 ggplot(titanic) +  
2   geom_boxplot(aes(x = class, y = age, fill = survived), alpha = 0.5)
```

Warning: Removed 263 rows containing non-finite values (`stat\_boxplot()`).



## 4 Smoking and Pregnancy



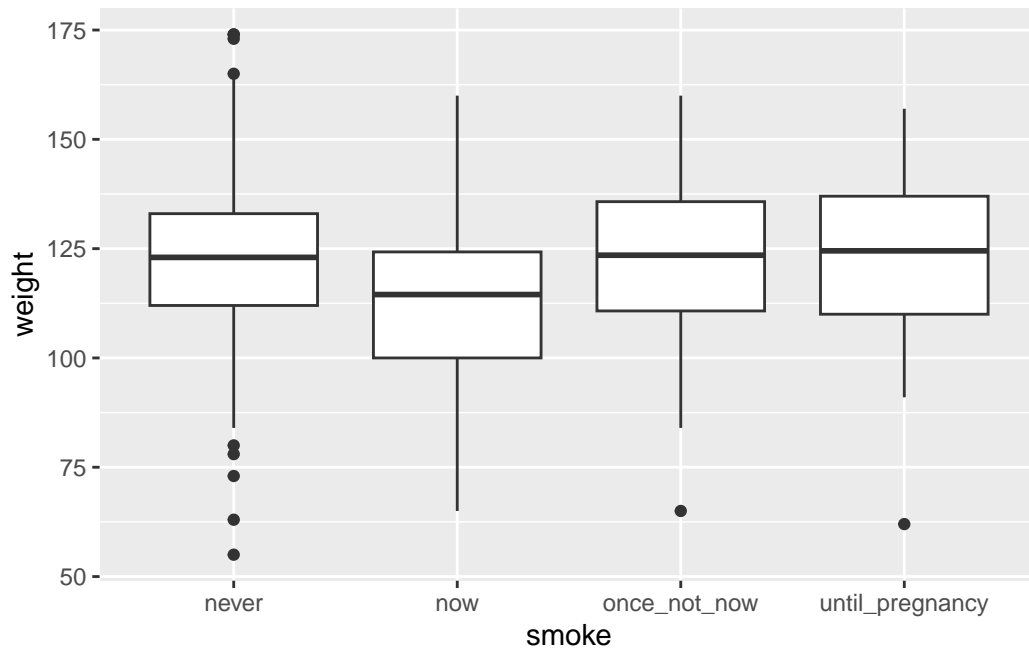
## 4.1 Load the dataset

```
1 smoking =  
  ↪ read.csv("https://raw.githubusercontent.com/ahmedmoustafa/datasets/main/smoking/smoking.csv")  
2 head(smoking)
```

id	dategestat	weightparity	commaceragenmedm	heightdweightblacel.alld.heightweightl.tosmoke	quit.totgss
15	141284	120	1 asian27	5 62 100 asian31	5 65 110 1 1 never 0 0
20	1499282	113	2 white33	5 64 135 white38	5 70 148 1 4 never 0 0
100	1673286	136	4 white25	2 62 93 white28	2 64 130 1 4 until_pregnancy
129	1562245	132	2 black23	1 65 140 black23	4 71 192 1 2 never 0 0
142	1402289	120	3 white25	4 62 125 white26	1 70 180 0 2 never 0 0
171	1593282	144	4 white32	2 64 124 white36	1 74 185 1 2 now 1 1

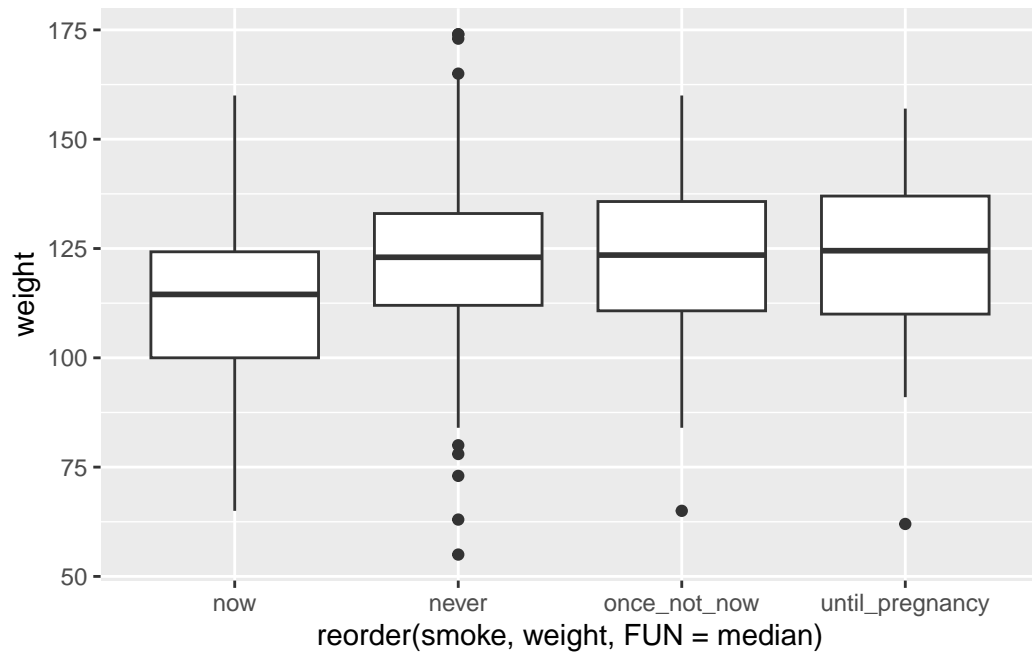
## 4.2 Mom's smoking and baby's weight

```
1 ggplot(smoking) +  
2   geom_boxplot(aes(x = smoke, y = weight))
```



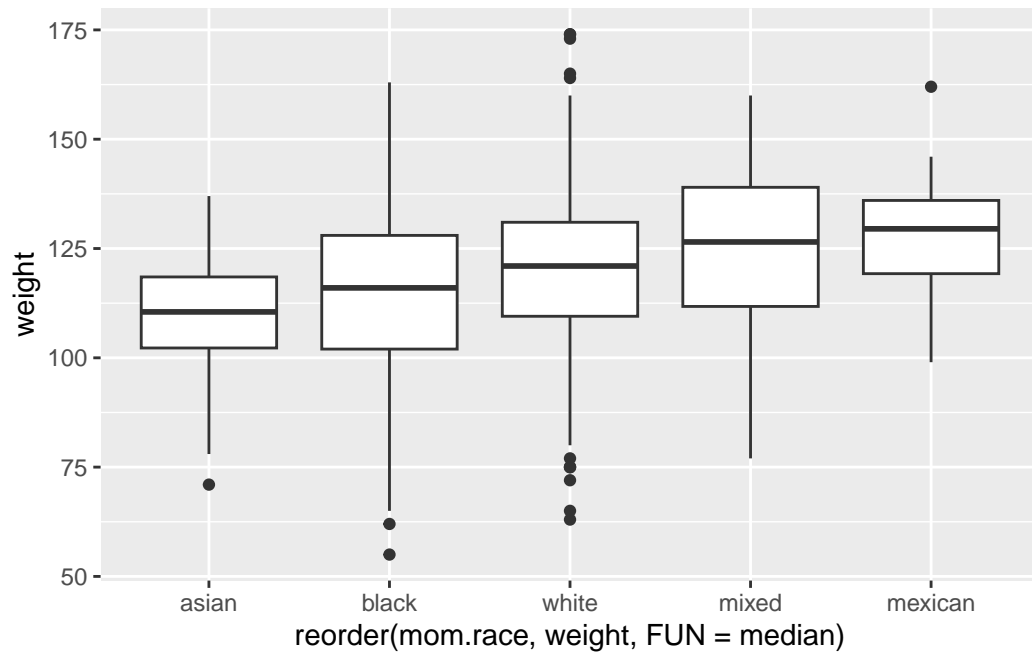
## 4.3 Mom's smoking and baby's weight with reordered x-axis

```
1 ggplot(smoking) +  
2   geom_boxplot(aes(x = reorder(smoke, weight, FUN = median), y =  
3     weight))
```



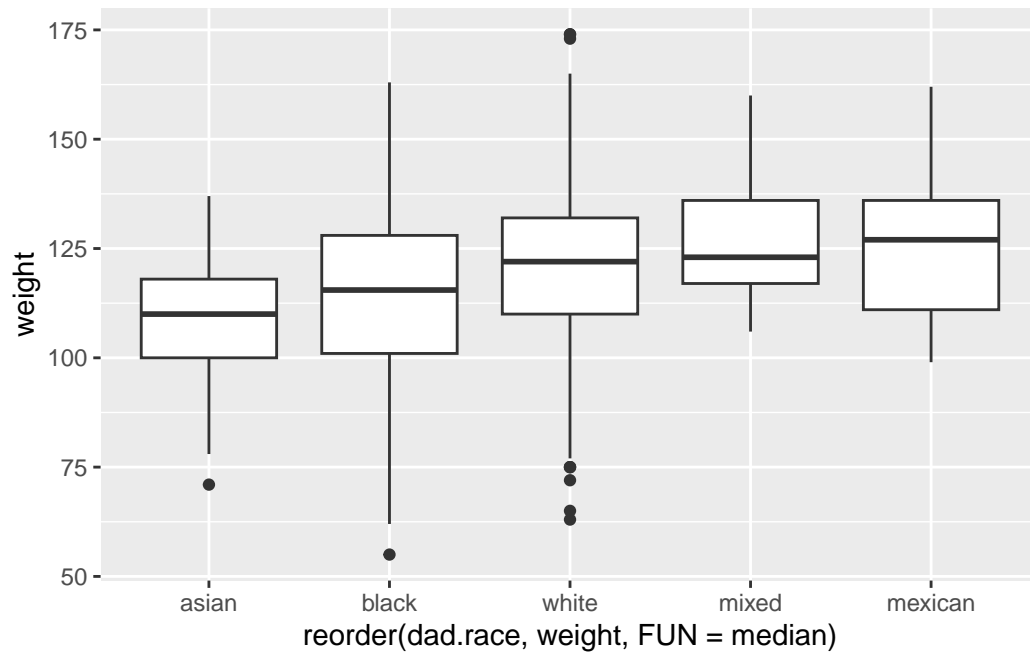
#### 4.4 Mom's race and baby's weight

```
1 ggplot(smoking) +  
2   geom_boxplot(aes(x = reorder(mom.race, weight, FUN = median), y =  
   ↪ weight))
```



#### 4.5 Dad's race and baby's weight

```
1 ggplot(smoking) +  
2   geom_boxplot(aes(x = reorder(dad.race, weight, FUN = median), y =  
   ↪ weight))
```



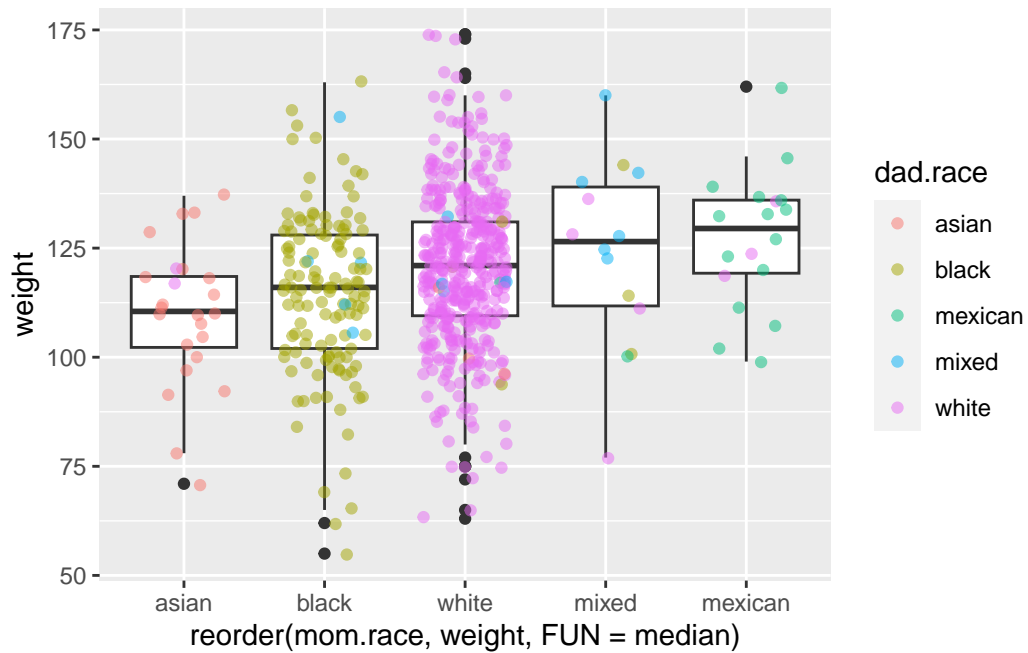
#### 4.6 Mom's race and baby's weight and dad's race

```

1 ggplot(smoking) +
2   geom_boxplot(aes(x = reorder(mom.race, weight, FUN = median), y =
   ↪ weight)) +
3   geom_jitter(aes(x = reorder(mom.race, weight, FUN = median), y =
   ↪ weight, color = dad.race), alpha = 0.5, width = 0.3)

```





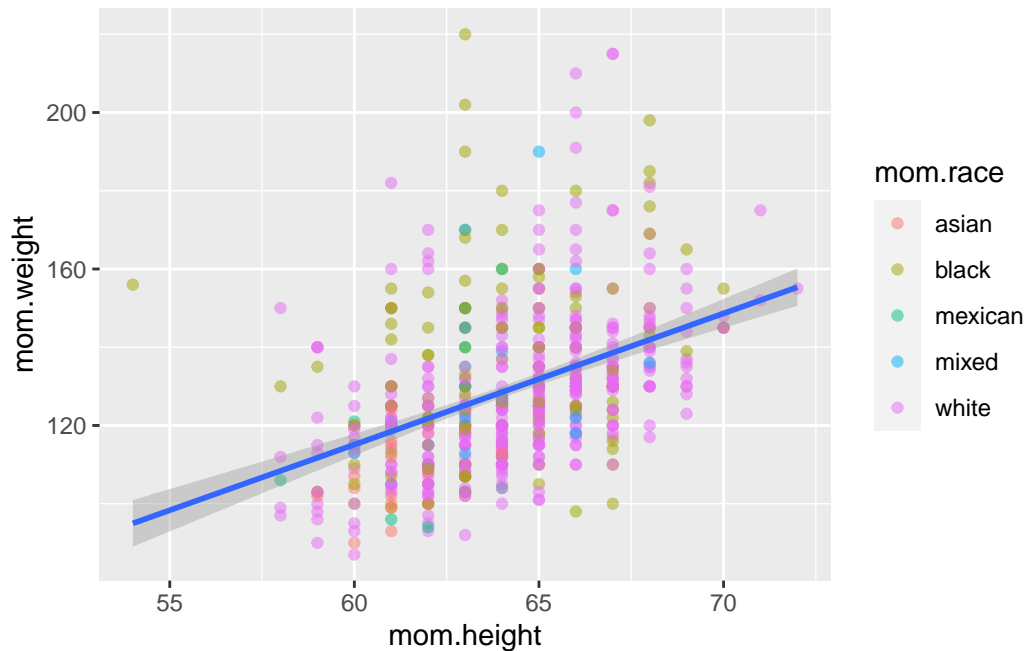
#### 4.7 Mom's height and moms's weight

```

1 ggplot(smoking) +
2   geom_point(aes(x = mom.height, y = mom.weight, color = mom.race),
3     ↪ alpha = 0.5) +
4   geom_smooth(aes(x = mom.height, y = mom.weight), method = "lm")

```

`geom\_smooth()` using formula = 'y ~ x'



```
1 model = lm (data = smoking, formula = mom.weight ~ mom.height)
2 summary(model)
```

Call:

```
lm(formula = mom.weight ~ mom.height, data = smoking)
```

Residuals:

Min	1Q	Median	3Q	Max
-38.579	-11.933	-3.515	7.276	94.839

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	-86.174	18.671	-4.615	4.79e-06 ***
mom.height	3.354	0.291	11.526	< 2e-16 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 18.55 on 608 degrees of freedom

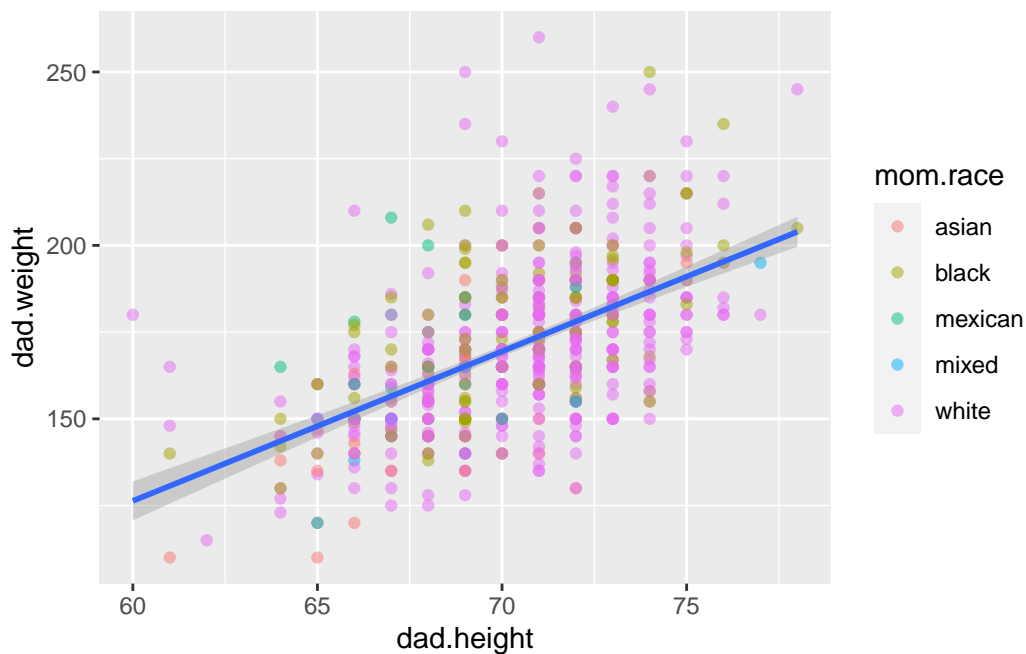
Multiple R-squared: 0.1793, Adjusted R-squared: 0.178

F-statistic: 132.8 on 1 and 608 DF, p-value: < 2.2e-16

## 4.8 Dad's height and dad's weight

```
1 ggplot(smoking) +  
2   geom_point(aes(x = dad.height, y = dad.weight, color = mom.race),  
3     ↪ alpha = 0.5) +  
4   geom_smooth(aes(x = dad.height, y = dad.weight), method = "lm")
```

`geom\_smooth()` using formula = 'y ~ x'



```
1 model = lm (data = smoking, formula = dad.weight ~ dad.height)  
2 summary(model)
```

Call:

```
lm(formula = dad.weight ~ dad.height, data = smoking)
```

Residuals:

Min	1Q	Median	3Q	Max
-48.067	-13.067	-1.825	10.554	86.243

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	-132.2898	18.8057	-7.035	5.4e-12	***
dad.height	4.3105	0.2674	16.120	< 2e-16	***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

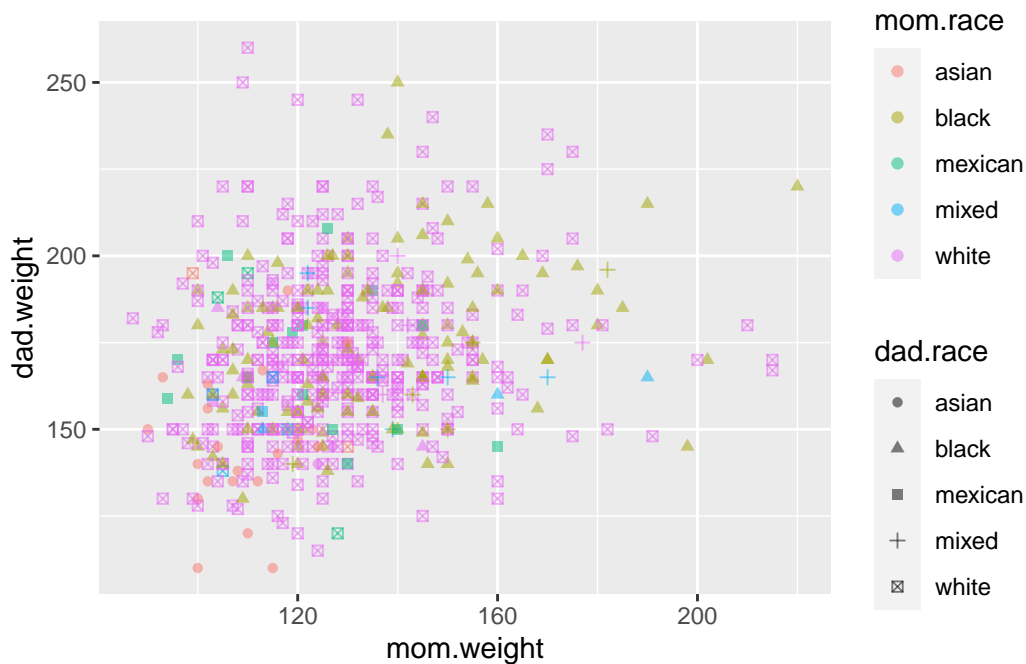
Residual standard error: 19.02 on 608 degrees of freedom

Multiple R-squared: 0.2994, Adjusted R-squared: 0.2983

F-statistic: 259.9 on 1 and 608 DF, p-value: < 2.2e-16

## 4.9 Mom's weight and dad's weight

```
1 ggplot(smoking) +  
2   geom_point(aes(x = mom.weight, y = dad.weight, color = mom.race, shape  
  ↪   = dad.race), alpha = 0.5)
```



```
1 model = lm (data = smoking, formula = dad.weight ~ mom.weight)  
2 summary(model)
```

Call:

```
lm(formula = dad.weight ~ mom.weight, data = smoking)
```

Residuals:

Min	1Q	Median	3Q	Max
-57.481	-15.817	-2.051	14.097	93.646

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )
(Intercept)	141.54810	5.74900	24.621	< 2e-16 ***
mom.weight	0.22551	0.04407	5.117	4.16e-07 ***

---

Signif. codes: 0 '\*\*\*' 0.001 '\*\*' 0.01 '\*' 0.05 '.' 0.1 ' ' 1

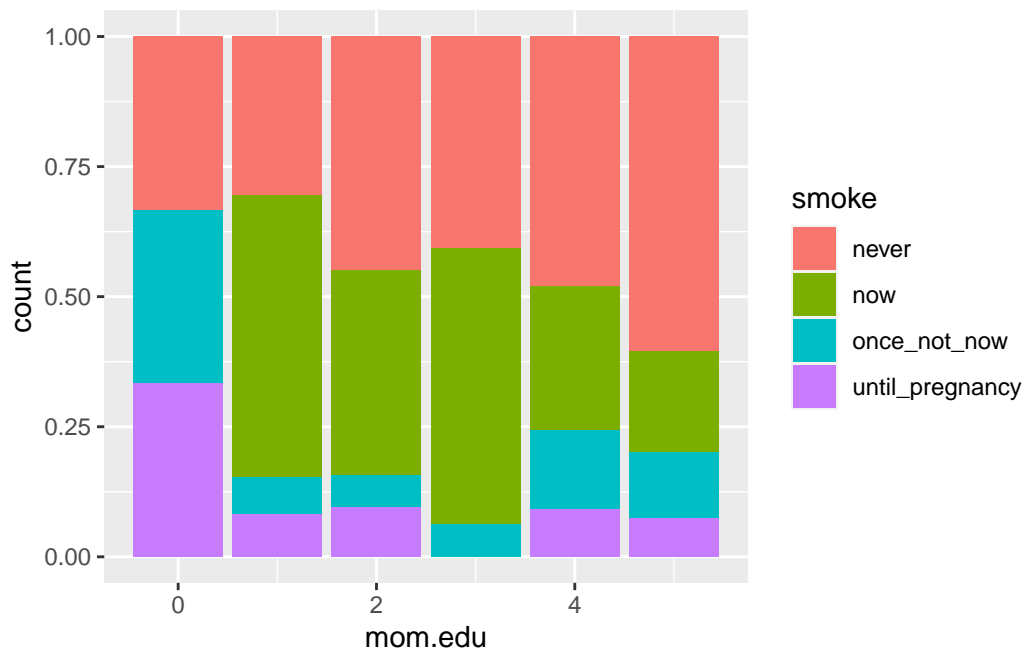
Residual standard error: 22.25 on 608 degrees of freedom

Multiple R-squared: 0.04129, Adjusted R-squared: 0.03972

F-statistic: 26.19 on 1 and 608 DF, p-value: 4.159e-07

#### 4.10 Mom's smoking and mom's education

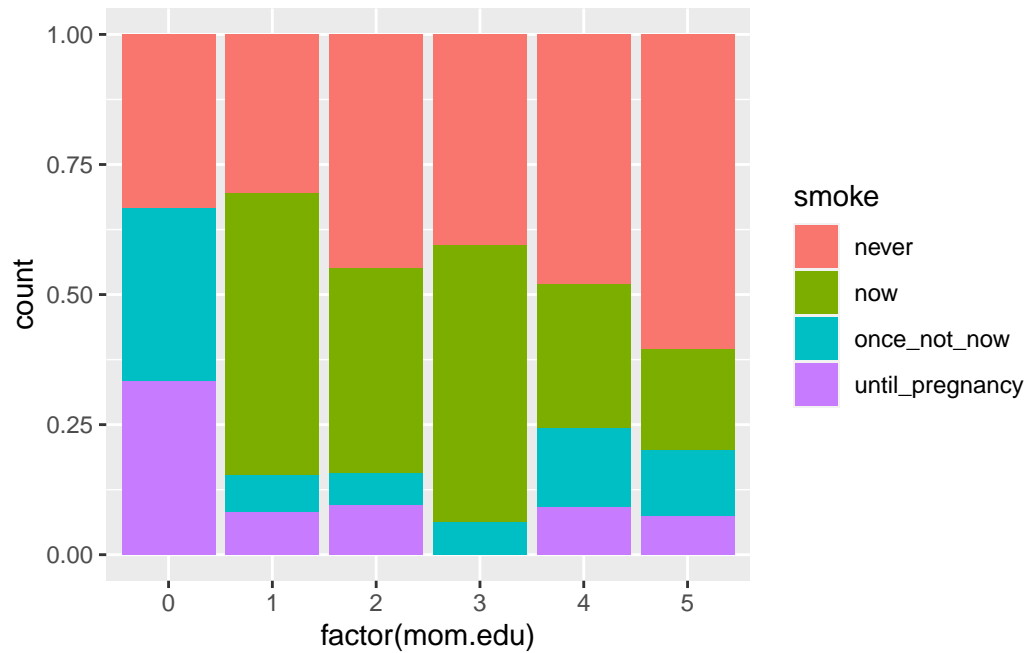
```
1 ggplot(smoking) +  
2   geom_bar(aes(x = mom.edu, fill = smoke), position = "fill")
```



```

1 ggplot(smoking) +
2   geom_bar(aes(x = factor(mom.edu), fill = smoke), position = "fill")

```

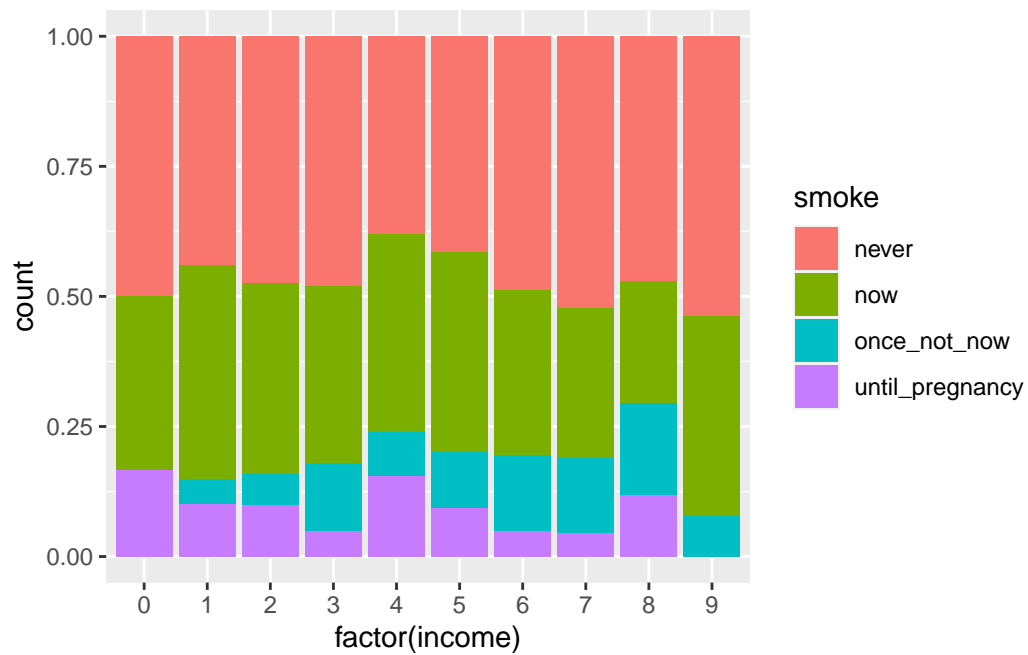


#### 4.11 Mom's smoking and the family's income

```

1 ggplot(smoking) +
2   geom_bar(aes(x = factor(income), fill = smoke), position = "fill")

```

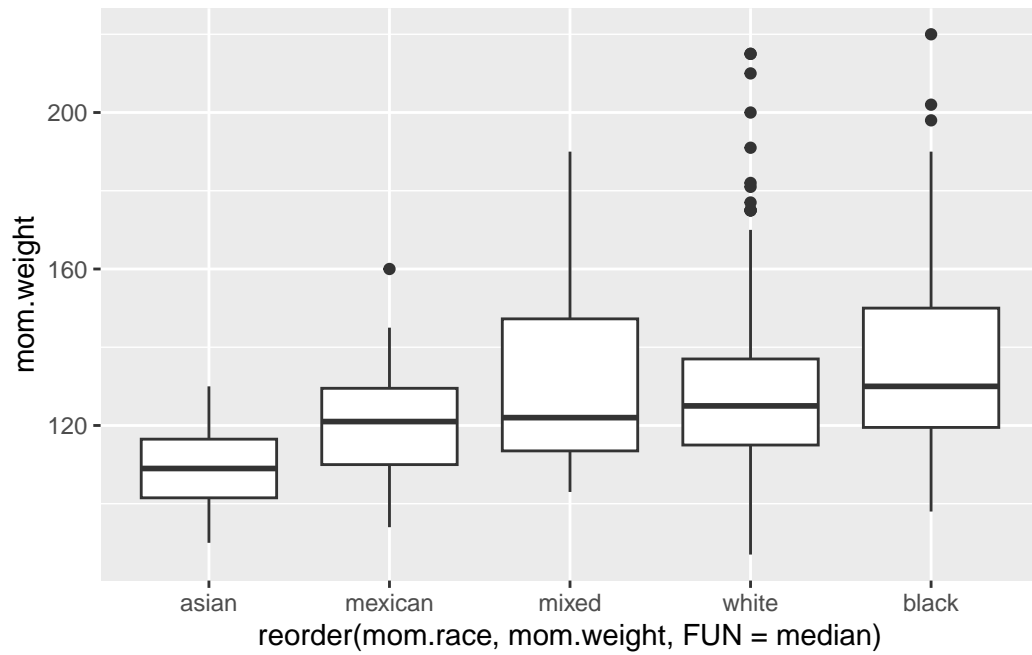


#### 4.12 Mom's race and mom's weight

```

1 ggplot(smoking) +
2   geom_boxplot(aes(x = reorder(mom.race, mom.weight, FUN = median), y =
   ↪ mom.weight))

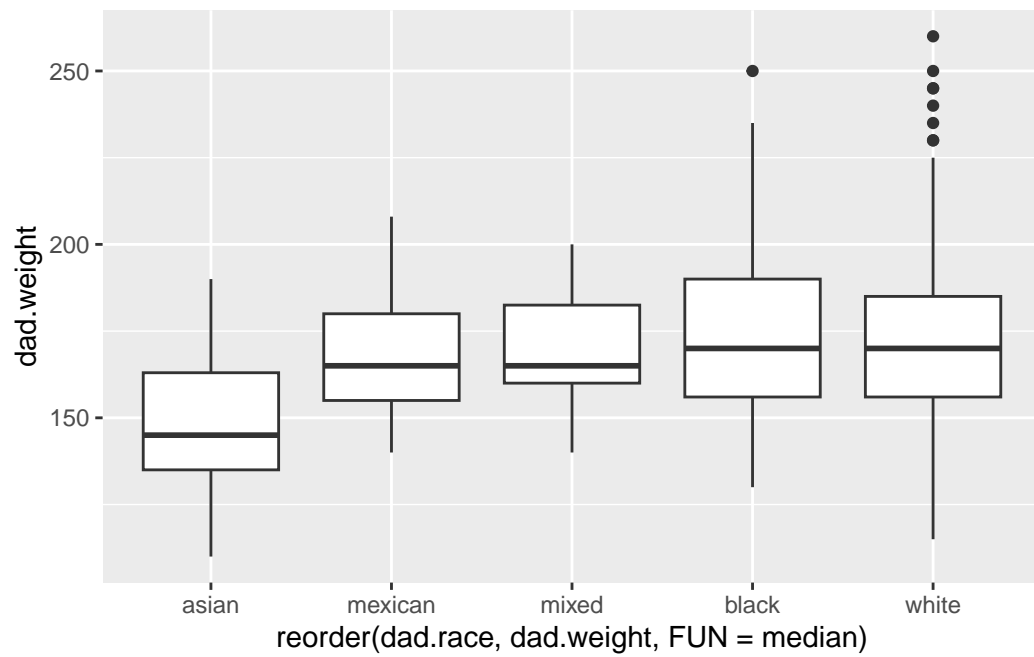
```



#### 4.13 Dad's race and dad's weight

```
1 ggplot(smoking) +  
2   geom_boxplot(aes(x = reorder(dad.race, dad.weight, FUN = median), y =  
   ↪ dad.weight))
```





## 5 The End

