

# Data manipulation with dplyr

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# DPLYR

*Basics*

# dplyr

❖ The new coup of **Hadley Wickham**, creator of

⇒ ggplot2

⇒ plyr

# plyr

- ❖ functional programming paradigm
- ❖ Functions as parameters of second level functions
  - ⇒ `lply(some.list, some.function)` # returns list
  - ⇒ `laply(some.list, some.function)` # returns array
  - ⇒ `aapply(some.array, some.function)` # returns array
  - ⇒ ...
- ❖ replaces classic second level functions: `apply`, `lapply`, `sapply`, `replicate`

# dplyr

next generation data.frame manipulation

- ❖ Simple interface
- ❖ Readable code
- ❖ Fast
- ❖ Can transparently deal with remote data
- ❖ interfaces well with → plyr and →ggplot

# Basic elements of plyr

## ❖ Functions:

- ⇒ Filter
- ⇒ Select
- ⇒ Mutate
- ⇒ Group\_by
- ⇒ Summarize
- ⇒ Arrange

## ❖ Operator

- ⇒ Concatenation by `%.>%` or `%>%>%`

# Example 1: filter some observations

Classic

```
D[S$subject == 4 & D$trial == 10,]
```

dplyr

```
filter(D, subject == 4, trial == 10)
```

## Example 2: select some variables

Classic

```
D[,c("subject", "trial")]
```

dplyr

```
select(D, subject, trial)
```



## Example 3: add a variable

Classic

```
D$freq <- D$count/D$time
```

dplyr

```
D <- mutate(D, freq = count/time)
```

# Example 4: summarize data

## Classic

```
aggregate(D$RT, list(subject =  
D$Subject), mean)
```

## dplyr

```
group_by(D, subject) %>%  
summarize(totalcount = sum(count))
```

# Example 5: order data

Classic

```
D[order(D$subject, D$trial),]
```

dplyr

```
arrange(D, subject, trial)
```



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Pipelining

# Piping commands with %.%

- ❖ `select(D, subject, trial, count, time) %.%  
 manipulate(freq = count/time) %.%  
 group_by(subject) %.%  
 summarize(avgfreq = mean(freq))`
- ❖ You can even:  
 `D %.% select(subject, trial, count, time) %.% ...`
- ❖ Or:  
 `read.spss("D.sav", to.data.frame = T) %.%  
 select(subject, trial, count, time)`

# Interfacing with ggplot

```
D%.%  
mutate(freq = count/time) %.%  
ggplot(aes(participant, freq))
```

This seems to work for all functions that take the data.frame as first argument

# Interfacing with lm

**lm(formula, data)** ← data.frame not first arg

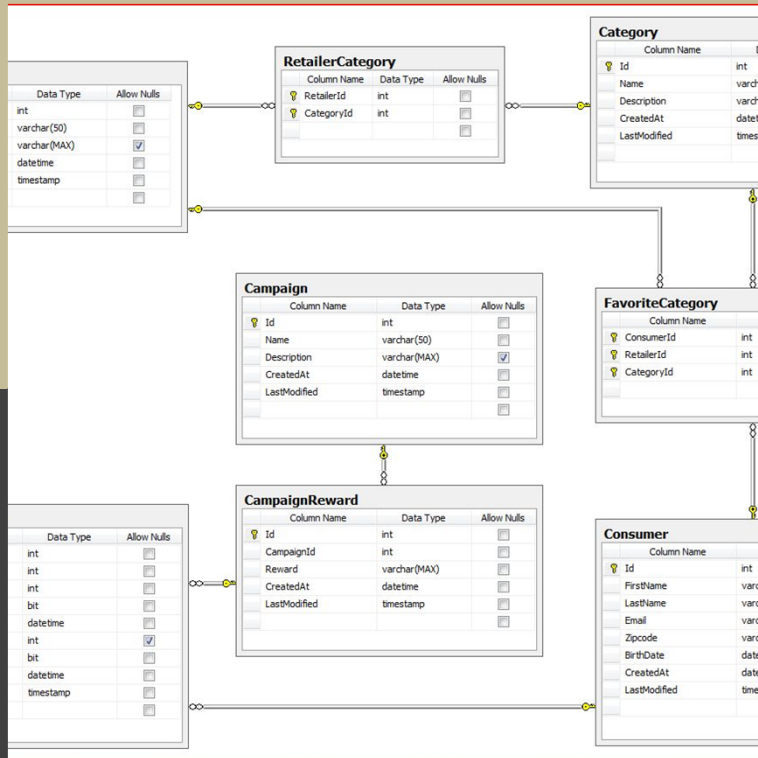
New in dplyr: `%>%` (magrittr)

`D%>%`

`mutate(freq = count/time) %>%`

`lm(freq ~ age, data = .) %>%`

`summary()`



# PRINCIPLES OF DATA MODELLING



# Tables, Keys, Redundancy

Subj	Gender	Word	Picture	RT
1	m	speak	robo 1	352
1	m	speak	robo 2	789
1	m	beep	robo 1	435
1	m	beep	robo 2	978
2	f	speak	robo 1	1423
2	f	speak	robo 2	1453
2	f	beep	robo 1	983
2	f	beep	robo 2	1234

- ❖ **Key:** smallest combination of variables that identifies an observation
- ❖ **Redundancy:** value of one column is strictly determined by another column

# Avoiding redundancy

Subj		Word	Picture	RT
1		speak	robo 1	352
1		speak	robo 2	789
1		beep	robo 1	435
1		beep	robo 2	978
2		speak	robo 1	1423
2		speak	robo 2	1453
2		beep	robo 1	983
2		beep	robo 2	1234

Subj		Gender
1		m
2		f

# re-joining

```
join(ExperimentalData,  
     SubjectData,  
     by = Subj)
```

*In data modeling speak, this is a  
1:n relation with  
Subj as foreign key*

Subj	Gender	Word	Picture	RT
1	m	speak	robo 1	352
1	m	speak	robo 2	789
1	m	beep	robo 1	435
1	m	beep	robo 2	978
2	f	speak	robo 1	1423
2	f	speak	robo 2	1453
2	f	beep	robo 1	983
2	f	beep	robo 2	1234

join commands supplied by plyr

# TIPPS & TRICKS

# plyr and dplyr

- ❖ Always load plyr first  
library(plyr)  
library(dplyr)
- ❖ If accidentally done wrong: Restart R
- ❖ plyr even warns you

# Renaming a column

## Classic

```
D <- read.spss("D.sav", to.data.frame = T)
```

```
D$participant <- D$proefpersoon
```

```
D$proefpersoon <- NULL
```

## dplyr

```
D <- read.spss("D.sav", to.data.frame = T) %>%
```

```
  select(participant = proefpersoon, trial:time)
```