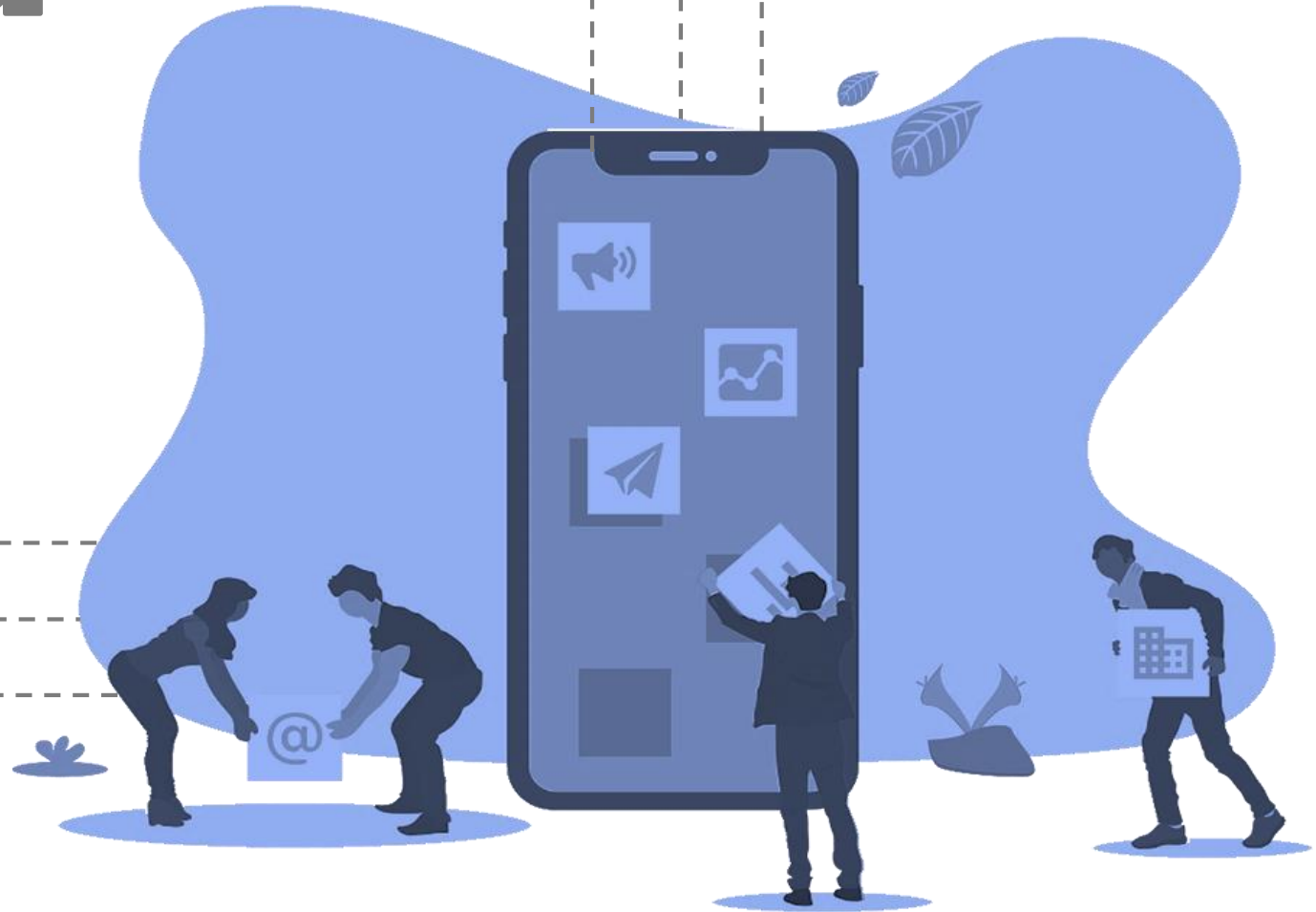


The Data Science Track



Prepared By: R. Daynolo

INTRODUCTION TO DATA SCIENCE WITH R



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PROCESS



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KNOWLEDGE

20. Managing and Reshaping of Data

CREATING NEW VARIABLES

Why create new variables?

- Often the raw data won't have a value you are looking for
- You will need to transform the data to get the values you would like
- Usually you will add those values to the data frames you are working with
- Common variables to create
 - Missingness indicators
 - “Cutting up” quantitative variables
 - Applying transformations



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COMMON TRANSFORMS

- **abs(x)** absolute value
- **sqrt(x)** square root
- **ceiling(x)** ceiling(3.475) is 4
- **floor(x)** floor(3.475) is 3
- **round(x, digits=n)** round(3.475, digits=2) is 3.48
- **signif(x, digits=n)** signif(3.475, digits=2) is 3.5
- **cos(x), sin(x)** etc
- **log(x)** natural logarithm
- **log2(x), log10(x)** other common logs
- **exp(x)** exponentiating x



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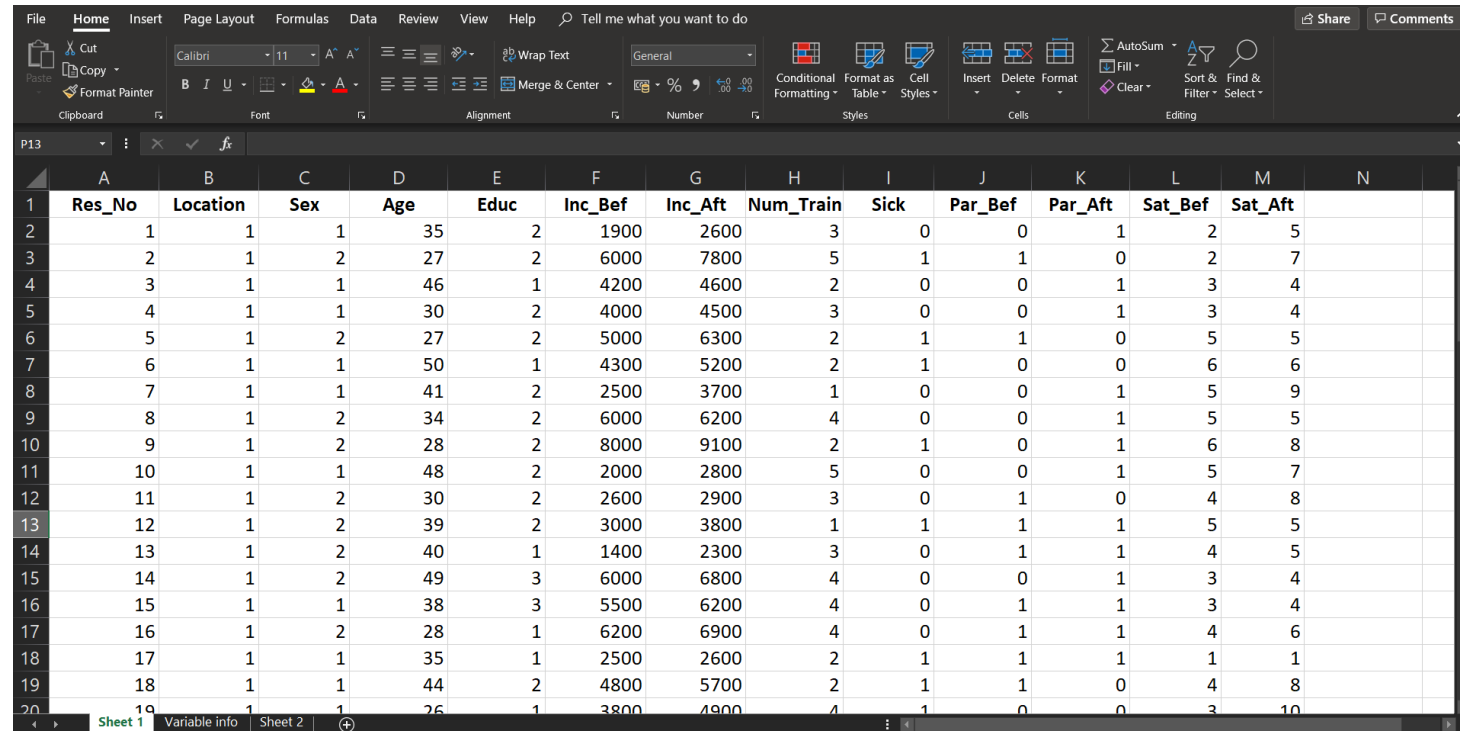
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RESHAPING DATA

The goal is tidy data



	A	B	C	D	E	F	G	H	I	J	K	L	M	N
1	Res_No	Location	Sex	Age	Educ	Inc_Bef	Inc_Aft	Num_Train	Sick	Par_Bef	Par_Aft	Sat_Bef	Sat_Aft	
2	1	1	1	35	2	1900	2600	3	0	0	1	2	5	
3	2	1	2	27	2	6000	7800	5	1	1	0	2	7	
4	3	1	1	46	1	4200	4600	2	0	0	1	3	4	
5	4	1	1	30	2	4000	4500	3	0	0	1	3	4	
6	5	1	2	27	2	5000	6300	2	1	1	0	5	5	
7	6	1	1	50	1	4300	5200	2	1	0	0	6	6	
8	7	1	1	41	2	2500	3700	1	0	0	1	5	9	
9	8	1	2	34	2	6000	6200	4	0	0	1	5	5	
10	9	1	2	28	2	8000	9100	2	1	0	1	6	8	
11	10	1	1	48	2	2000	2800	5	0	0	1	5	7	
12	11	1	2	30	2	2600	2900	3	0	1	0	4	8	
13	12	1	2	39	2	3000	3800	1	1	1	1	5	5	
14	13	1	2	40	1	1400	2300	3	0	1	1	4	5	
15	14	1	2	49	3	6000	6800	4	0	0	1	3	4	
16	15	1	1	38	3	5500	6200	4	0	1	1	3	4	
17	16	1	2	28	1	6200	6900	4	0	1	1	4	6	
18	17	1	1	35	1	2500	2600	2	1	1	1	1	1	
19	18	1	1	44	2	4800	5700	2	1	1	0	4	8	
20	19	1	1	26	1	3800	4900	4	1	0	0	3	10	

1. Each variable forms a column
2. Each observation forms a row
3. Each table/file stores data about one kind of observation (e.g. people)



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START WITH RESHAPING

```
> library(reshape2)
> head(mtcars)
```

	mpg	cyl	disp	hp	drat	wt	qsec	vs	am	gear	carb
Mazda RX4	21.0	6	160	110	3.90	2.620	16.46	0	1	4	4
Mazda RX4 Wag	21.0	6	160	110	3.90	2.875	17.02	0	1	4	4
Datsun 710	22.8	4	108	93	3.85	2.320	18.61	1	1	4	1
Hornet 4 Drive	21.4	6	258	110	3.08	3.215	19.44	1	0	3	1
Hornet Sportabout	18.7	8	360	175	3.15	3.440	17.02	0	0	3	2
Valiant	18.1	6	225	105	2.76	3.460	20.22	1	0	3	1



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MELTING DATA FRAMES

```
mtcars$carname <- rownames(mtcars)
carMelt <- melt(mtcars, id=c("carname", "gear", "cyl"),
               measure.vars = c("mpg", "hp"))
```

```
> head(carMelt, n = 3)
      carname gear cyl variable value
1   Mazda RX4   4   6      mpg    21.0
2 Mazda RX4 Wag   4   6      mpg    21.0
3  Datsun 710    4   4      mpg    22.8
>
> tail(carMelt, n = 3)
      carname gear cyl variable value
62 Ferrari Dino   5   6      hp    175
63 Maserati Bora   5   8      hp    335
64  Volvo 142E    4   4      hp    109
```



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CASTING DATA FRAMES

```
> cylData <- dcast(carMelt, cyl ~ variable)
Aggregation function missing: defaulting to length
```

```
> cylData
  cyl mpg hp
1   4  11 11
2   6   7  7
3   8  14 14
```

```
> cylData <- dcast(carMelt, cyl ~ variable, mean)
```

```
> cylData
  cyl      mpg      hp
1   4 26.66364 82.63636
2   6 19.74286 122.28571
3   8 15.10000 209.21429
```



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AVERAGING VALUES

```
> head(InsectSprays)
```

	count	spray
1	10	A
2	7	A
3	20	A
4	14	A
5	14	A
6	12	A

```
> tapply(InsectSprays$count, InsectSprays$spray, sum)
```

A	B	C	D	E	F
174	184	25	59	42	200



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ANOTHER WAY - SPLIT

```
> spIns <- split(InsectSprays$count, InsectSprays$spray)
> spIns
$A
 [1] 10  7 20 14 14 12 10 23 17 20 14
[12] 13

$B
 [1] 11 17 21 11 16 14 17 17 19 21  7
[12] 13

$C
 [1] 0 1 7 2 3 1 2 1 3 0 1 4

$D
 [1]  3  5 12  6  4  3  5  5  5  5  2
[12]  4

$E
```

ANOTHER WAY - APPLY

```
> sprCount <- lapply(spIns, sum)
> sprCount
$A
[1] 174

$B
[1] 184

$C
[1] 25

$D
[1] 59

$E
[1] 12
```



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ANOTHER WAY - COMBINE

```
> unlist(sprCount)
```

A	B	C	D	E	F
174	184	25	59	42	200

```
> sapply(spIns, sum)
```

A	B	C	D	E	F
174	184	25	59	42	200



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ANOTHER WAY – PLYR PACKAGE

```
> library(plyr)
> ddply(InsectSprays, .(spray), summarize, sum = sum(count))
```

	spray	sum
1	A	174
2	B	184
3	C	25
4	D	59
5	E	42
6	F	200



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CREATING A NEW VARIABLE

```
spraySums <- ddpIy(InsectSprays, .(spray), summarize,  
                    sum = ave(count, FUN=sum))
```

```
> dim(spraySums)  
[1] 72  2  
> head(spraySums)  
  spray sum  
1     A 174  
2     A 174  
3     A 174  
4     A 174  
5     A 174  
6     A 174
```



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MORE INFORMATION

- A tutorial from the developer of plyr
<http://plyr.had.co.nz/09-user/>
- A nice reshape tutorial
<https://www.slideshare.net/jeffreybreen/reshaping-data-in-r>
- A good plyr primer
<https://www.r-bloggers.com/a-quick-primer-on-split-apply-combine-problems/>
- See also the functions
 - acast – for casting as multi-dimensional arrays
 - arrange – for faster reordering without using order() commands
 - mutate – adding new variables



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MANAGING DATA FRAMES WITH DPLYR

dplyr

The data frame is a key data structure in statistics and in R

- There is one observation per row
- Each column represents a variable or measure or characteristic
- Primary implementation that you will use is the default R implementation
- Other implementations, particularly relational databases systems



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MANAGING DATA FRAMES WITH DPLYR

dplyr

- Developed by Hadley Wickham of Rstudio
- An optimized and distilled version of plyr package (also by Hadley)
- Does not provide any “new” functionality per se, but greatly simplifies existing functionality in R
- Provides a “grammar” (in particular, verbs) for data manipulation
- Is very fast, as many key operations are coded in C++



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DPLYR VERBS

- `select`: return a subset of the columns of the data frame
- `filter`: extract a subset of rows from a data frame based on logical conditions
- `arrange`: reorder rows of a data frame
- `rename`: rename variables in a data frame
- `mutate`: add new variables/columns or transform existing variables
- `summarise` / `summarize`: generate summary statistics of different variables in the data frame, possibly within strata

There is also a handy `print` method that prevents you from printing a lot of data to the console.



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DPLYR PROPERTIES

- The first argument is a data frame.
- The subsequent arguments describe what to do with it, and you can refer to columns in the data frame directly without using the \$ operator (just use the names).
- The result is a new data frame
- Data frames must be properly formatted and annotated for this to all be useful



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SELECT

```
> chicago <- readRDS("chicago.rds")
> dim(chicago)
[1] 6940      8
> head(select(chicago, 1:5))
  city tmpd   dptp   date pm25tmean2
1 chic 31.5 31.500 1987-01-01        NA
2 chic 33.0 29.875 1987-01-02        NA
3 chic 33.0 27.375 1987-01-03        NA
4 chic 29.0 28.625 1987-01-04        NA
5 chic 32.0 28.875 1987-01-05        NA
6 chic 40.0 35.125 1987-01-06        NA
```



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SELECT

```
> names(chicago)[1:3]
[1] "city" "tmpd" "dptp"
> head(select(chicago, city:dptp))
  city tmpd  dptp
1 chic 31.5 31.500
2 chic 33.0 29.875
3 chic 33.0 27.375
4 chic 29.0 28.625
5 chic 32.0 28.875
6 chic 40.0 35.125
```



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SELECT

In dplyr you can do

```
> head(select(chicago, -(city:dptp)))
```

	date	pm25tmean2	pm10tmean2	o3tmean2	no2tmean2
1	1987-01-01	NA	34.00000	4.250000	19.98810
2	1987-01-02	NA	NA	3.304348	23.19099
3	1987-01-03	NA	34.16667	3.333333	23.81548
4	1987-01-04	NA	47.00000	4.375000	30.43452
5	1987-01-05	NA	NA	4.750000	30.33333
6	1987-01-06	NA	48.00000	5.833333	25.77233



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SELECT

Equivalent base R

```
> i <- match("city", names(chicago))
> j <- match("dptp", names(chicago))
> head(chicago[, -(i:j)])
```

	date	pm25tmean2	pm10tmean2	o3tmean2	no2tmean2
1	1987-01-01	NA	34.00000	4.250000	19.98810
2	1987-01-02	NA	NA	3.304348	23.19099
3	1987-01-03	NA	34.16667	3.333333	23.81548
4	1987-01-04	NA	47.00000	4.375000	30.43452
5	1987-01-05	NA	NA	4.750000	30.33333
6	1987-01-06	NA	48.00000	5.833333	25.77233



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FILTER

```
> chicago.f <- filter(chicago, pm25tmean2>30)
> head(select(chicago.f, 1:3, pm25tmean2), 10)
```

	city	tmpd	dptp	pm25tmean2
1	chic	23	21.9	38.10
2	chic	28	25.8	33.95
3	chic	55	51.3	39.40
4	chic	59	53.7	35.40
5	chic	57	52.0	33.30
6	chic	57	56.0	32.10
7	chic	75	65.8	56.50
8	chic	61	59.0	33.80
9	chic	73	60.3	30.30
10	chic	78	67.1	41.40



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FILTER

```
> chicago.f <- filter(chicago, pm25tmean2>30 & tmpd>80)  
> head(select(chicago.f, 1:3, pm25tmean2),10)
```

	city	tmpd	dptp	pm25tmean2
1	chic	81	71.2	39.6000
2	chic	81	70.4	31.5000
3	chic	82	72.2	32.3000
4	chic	84	72.9	43.7000
5	chic	85	72.6	38.8375
6	chic	84	72.6	38.2000
7	chic	82	67.4	33.0000
8	chic	82	63.5	42.5000
9	chic	81	70.4	33.1000
10	chic	82	66.2	38.8500



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ARRANGE

Reordering rows of a data frame (while preserving corresponding order of other columns) is normally a pain to do in R.

```
> chicago <- arrange(chicago, date)
> head(select(chicago, date, pm25tmean2), 3)
      date pm25tmean2
1 1987-01-01      NA
2 1987-01-02      NA
3 1987-01-03      NA
> tail(select(chicago, date, pm25tmean2), 3)
      date pm25tmean2
6938 2005-12-29    7.45000
6939 2005-12-30   15.05714
6940 2005-12-31   15.00000
```



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ARRANGE

Columns can be arranged in descending order too.

```
> chicago <- arrange(chicago, desc(date))
> head(select(chicago, date, pm25tmean2), 3)
      date pm25tmean2
1 2005-12-31    15.00000
2 2005-12-30    15.05714
3 2005-12-29     7.45000
> tail(select(chicago, date, pm25tmean2), 3)
      date pm25tmean2
6938 1987-01-03      NA
6939 1987-01-02      NA
6940 1987-01-01      NA
```



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RENAME

Renaming a variable in a data frame in R is surprisingly hard to do!

```
> head(chicago[, 1:5], 3)
  city tmpd dptp      date pm25tmean2
1 chic   35 30.1 2005-12-31   15.00000
2 chic   36 31.0 2005-12-30   15.05714
3 chic   35 29.4 2005-12-29    7.45000
> chicago <- rename(chicago, dewpoint = dptp,
+                    pm25 = pm25tmean2)
> head(chicago[, 1:5], 3)
  city tmpd dewpoint      date      pm25
1 chic   35    30.1 2005-12-31 15.00000
2 chic   36    31.0 2005-12-30 15.05714
3 chic   35    29.4 2005-12-29  7.45000
```



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MUTATE

```
> chicago <- mutate(chicago, pm25detrend = pm25 - mean(pm25, na.rm=TRUE))  
> head(select(chicago, pm25, pm25detrend))
```

	pm25	pm25detrend
1	15.00000	-1.230958
2	15.05714	-1.173815
3	7.45000	-8.780958
4	17.75000	1.519042
5	23.56000	7.329042
6	8.40000	-7.830958



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GROUP_BY

Generating summary statistics by stratum

```
chicago <- mutate(chicago,  
                    tempcat = factor(1 * (tmpd > 80),  
                                     labels = c("cold", "hot")))  
hotcold <- group_by(chicago, tempcat)  
summarize(hotcold, pm25 = mean(pm25, na.rm = TRUE),  
           o3 = max(o3tmean2),  
           no2 = median(no2tmean2))
```

```
# A tibble: 3 x 4  
  tempcat    pm25    o3    no2  
  <fct>    <dbl> <dbl> <dbl>  
1 cold    15.978 66.588 24.549  
2 hot     26.481 62.970 24.939  
3 NA      47.738  9.4167 37.444
```



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GROUP_BY

```
chicago <- mutate(chicago,  
                    year = as.POSIXlt(date)$year + 1900)  
years <- group_by(chicago, year)  
summarize(years, pm25 = mean(pm25, na.rm = TRUE),  
           o3 = max(o3tmean2, na.rm = TRUE),  
           no2 = median(no2tmean2, na.rm = TRUE))
```

```
# A tibble: 19 x 4
```

	year <dbl>	pm25 <dbl>	o3 <dbl>	no2 <dbl>
1	<u>1987</u>	NaN	62.970	23.494
2	<u>1988</u>	NaN	61.677	24.523
3	<u>1989</u>	NaN	59.727	26.141
4	<u>1990</u>	NaN	52.229	22.596
5	<u>1991</u>	NaN	63.104	21.382
6	<u>1992</u>	NaN	50.829	24.789

%>%

```
chicago %>% mutate(month = as.POSIXlt(date)$mon + 1) %>%  
  group_by(month) %>%  
  summarize(pm25 = mean(pm25, na.rm = TRUE),  
            o3 = max(o3tmean2, na.rm = TRUE),  
            no2 = median(no2tmean2, na.rm = TRUE))
```

```
# A tibble: 12 x 4
```

	month	pm25	o3	no2
	<dbl>	<dbl>	<dbl>	<dbl>
1	1	17.770	28.222	25.354
2	2	20.375	37.375	26.780
3	3	17.408	39.050	26.770
4	4	13.859	47.949	25.031
5	5	14.074	52.75	24.222
6	6	15.865	66.588	25.011

MERGING DATA


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RESEARCH ARTICLE

Cooperation between Referees and Authors Increases Peer Review Accuracy


Jeffrey T. Leek , Margaret A. Taub, Fernando J. Pineda

Published: November 9, 2011 • <https://doi.org/10.1371/journal.pone.0026895>

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Abstract

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Abstract

Peer review is fundamentally a cooperative process between scientists in a community who agree to review each other's work in an unbiased fashion. Peer review is the foundation for decisions concerning publication in journals, awarding of grants, and academic promotion. Here we perform a laboratory study of open and closed peer review based on an online game. We show that when reviewer behavior was made public under open review, reviewers were rewarded for refereeing and formed significantly more cooperative interactions (13% increase in cooperation, $P=0.018$). We also show that referees and authors who participated in cooperative interactions had an 11% higher reviewing accuracy rate ($P=0.016$). Our results suggest that increasing cooperation in the peer review process can lead to a decreased risk of reviewing errors.

Figures

<https://journals.plos.org/plosone/article?id=10.1371/journal.pone.0026895>



MERGING DATA

```
> head(reviews)
```

	id	solution_id	reviewer_id	start	stop	time_left	accept
1	1	3	27	1304095698	1304095758	1754	1
2	2	4	22	1304095188	1304095206	2306	1
3	3	5	28	1304095276	1304095320	2192	1
4	4	1	26	1304095267	1304095423	2089	1
5	5	10	29	1304095456	1304095469	2043	1
6	6	2	29	1304095471	1304095513	1999	1

```
> head(solutions)
```

	id	problem_id	subject_id	start	stop	time_left	answer
1	1	156	29	1304095119	1304095169	2343	B
2	2	269	25	1304095119	1304095183	2329	C
3	3	34	22	1304095127	1304095146	2366	C
4	4	19	23	1304095127	1304095150	2362	D
5	5	605	26	1304095127	1304095167	2345	A
6	6	384	27	1304095131	1304095270	2242	C



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MERGING DATA – merge()

- Merges data frames
- Important parameter: x, y, by, by.x, by.y, all

```
> names(reviews)
[1] "id"          "solution_id" "reviewer_id" "start"
[5] "stop"        "time_left"   "accept"
> names(solutions)
[1] "id"          "problem_id"  "subject_id"  "start"
[5] "stop"        "time_left"   "answer"
```



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MERGING DATA – merge()

```
mergedData <- merge(reviews, solutions, by.x = "solution_id",  
                    by.y = "id", all = TRUE)
```

```
> head(mergedData)
```

	solution_id	id	reviewer_id	start.x	stop.x	time_left.x	accept
1	1	4	26	1304095267	1304095423	2089	1
2	2	6	29	1304095471	1304095513	1999	1
3	3	1	27	1304095698	1304095758	1754	1
4	4	2	22	1304095188	1304095206	2306	1
5	5	3	28	1304095276	1304095320	2192	1
6	6	16	22	1304095303	1304095471	2041	1

	problem_id	subject_id	start.y	stop.y	time_left.y	answer
1	156	29	1304095119	1304095169	2343	B
2	269	25	1304095119	1304095183	2329	C
3	34	22	1304095127	1304095146	2366	C
4	19	23	1304095127	1304095150	2362	D
5	605	26	1304095127	1304095167	2345	A
6	384	27	1304095131	1304095270	2242	C

DEFAULT – MERGE ALL COMMON COLUMN NAMES

```
> intersect(names(solutions), names(reviews))  
[1] "id"      "start"   "stop"    "time_left"
```

```
> mergedData2 <- merge(reviews, solutions, all = TRUE)
```

```
> head(mergedData2)
```

	id	start	stop	time_left	solution_id	reviewer_id	accept
1	1	1304095119	1304095169	2343	NA	NA	NA
2	1	1304095698	1304095758	1754	3	27	1
3	2	1304095119	1304095183	2329	NA	NA	NA
4	2	1304095188	1304095206	2306	4	22	1
5	3	1304095127	1304095146	2366	NA	NA	NA
6	3	1304095276	1304095320	2192	5	28	1

	problem_id	subject_id	answer
1	156	29	B
2	NA	NA	<NA>
3	269	25	C
4	NA	NA	<NA>
5	34	22	C
6	NA	NA	<NA>

USING JOIN IN THE PLYR PACKAGE

- Faster, but less full featured – defaults to left join, see help file for more

```
library(plyr)
df1 <- data.frame(id = sample(1:10), x = rnorm(10))
df2 <- data.frame(id = sample(1:10), y = rnorm(10))
```



ANALYSIS



STRUCTURE



ALGORITHM



PROCESS



PROGRAMMING



SOLVING



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USING JOIN IN THE PLYR PACKAGE

```
> arrange(join(df1, df2), id)
Joining by: id
```

	id	x	y
1	1	0.627027364	-0.60073051
2	2	0.700967679	0.03386791
3	3	-0.810659859	0.60660141
4	4	0.005873449	-1.31129057
5	5	0.380967414	0.73563731
6	6	-0.170144848	1.60118227
7	7	-1.698512322	-0.14455617
8	8	-0.110217865	-0.96965285
9	9	0.077961005	1.65329231
10	10	0.724992375	-1.94993964



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IF YOU HAVE MULTIPLE DATA FRAMES

```
df1 <- data.frame(id = sample(1:10), x = rnorm(10))  
df2 <- data.frame(id = sample(1:10), y = rnorm(10))  
df3 <- data.frame(id = sample(1:10), z = rnorm(10))  
dfList <- list(df1, df2, df3)
```



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IF YOU HAVE MULTIPLE DATA FRAMES

```
> join_all(dfList)
```

```
Joining by: id
```

```
Joining by: id
```

	id	x	y	z
1	6	1.3859707	-2.3950090	1.45381210
2	4	-0.5372697	2.2485589	0.58109302
3	10	-0.3127759	0.4397997	-0.01423728
4	7	-1.0128262	-0.1101474	-0.35719004
5	1	-0.5174372	0.1965517	0.67689562
6	9	0.5188877	-0.5958733	-0.02668022
7	5	-1.0822557	-2.3850438	-0.23531571
8	8	-0.3802740	0.2729542	-0.40372999
9	3	0.6611610	0.4494388	0.66859383
10	2	0.3689558	-0.8623506	-0.38679770



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MORE ON MERGING DATA

- The quick R data merging page
<https://www.statmethods.net/management/merging.html>
- plyr information
<http://plyr.had.co.nz/>



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