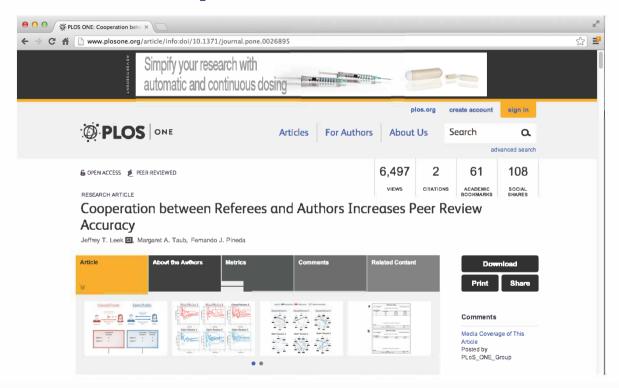


Merging data

Jeffrey Leek Johns Hopkins Bloomberg School of Public Health

Peer review experiment data



http://www.plosone.org/article/info:doi/10.1371/journal.pone.0026895

Peer review data

```
if(!file.exists("./data")){dir.create("./data")}
fileUrl1 = "https://dl.dropboxusercontent.com/u/7710864/data/reviews-apr29.csv"
fileUrl2 = "https://dl.dropboxusercontent.com/u/7710864/data/solutions-apr29.csv"
download.file(fileUrl1,destfile="./data/reviews.csv",method="curl")
download.file(fileUrl2,destfile="./data/solutions.csv",method="curl")
reviews = read.csv("./data/reviews.csv"); solutions <- read.csv("./data/solutions.csv")
head(reviews,2)</pre>
```

```
head(solutions,2)
```

Merging data - merge()

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

Merging data - merge()

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

```
solution id id reviewer id
                                           stop.x time left.x accept problem id subject id
                               start.x
                         26 1304095267 1304095423
                                                          2089
                                                                             156
                                                                                         29
                         29 1304095471 1304095513
                                                          1999
                                                                             269
                                                                                         25
                                                                                         22
                             1304095698 1304095758
                                                          1754
                                                                              34
                         22 1304095188 1304095206
                                                          2306
                                                                             19
                                                                                         23
                         28 1304095276 1304095320
                                                          2192
                                                                             605
                                                                                         26
           6 16
                          22 1304095303 1304095471
                                                                             384
                                                                                         27
                                                          2041
                stop.y time left.y answer
     start.y
1 1304095119 1304095169
                               2343
                                         В
                              2329
2 1304095119 1304095183
3 1304095127 1304095146
                              2366
4 1304095127 1304095150
                              2362
                                         D
5 1304095127 1304095167
                              2345
6 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	problem_id	subject_id	answer
1	1	1304095119	1304095169	2343	NA	NA	NA	156	29	В
2	1	1304095698	1304095758	1754	3	27	1	NA	NA	<na></na>
3	2	1304095119	1304095183	2329	NA	NA	NA	269	25	С
4	2	1304095188	1304095206	2306	4	22	1	NA	NA	<na></na>
5	3	1304095127	1304095146	2366	NA	NA	NA	34	22	С
6	3	1304095276	1304095320	2192	5	28	1	NA	NA	<na></na>

Using join in the plyr package

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

```
df1 = data.frame(id=sample(1:10),x=rnorm(10))
df2 = data.frame(id=sample(1:10),y=rnorm(10))
arrange(join(df1,df2),id)
```

```
id x y

1 1 0.2514 0.2286

2 2 0.1048 0.8395

3 3 -0.1230 -1.1165

4 4 1.5057 -0.1121

5 5 -0.2505 1.2124

6 6 0.4699 -1.6038

7 7 0.4627 -0.8060

8 8 -1.2629 -1.2848

9 9 -0.9258 -0.8276

10 10 2.8065 0.5794
```

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)
join_all(dfList)
```

```
id x y z

1 6 0.39093 -0.16670 0.56523

2 1 -1.90467 0.43811 -0.37449

3 7 -1.48798 -0.85497 -0.69209

4 10 -2.59440 0.39591 -0.36134

5 3 -0.08539 0.08053 1.01247

6 4 -1.63165 -0.13158 0.21927

7 5 -0.50594 0.24256 -0.44003

8 9 -0.85062 -2.08066 -0.96950

9 2 -0.63767 -0.10069 0.09002

10 8 1.20439 1.29138 -0.88586
```

More on merging data

- The quick R data merging page http://www.statmethods.net/management/merging.html
- plyr information http://plyr.had.co.nz/
- Types of joins http://en.wikipedia.org/wiki/Join_(SQL))