

3.0 Reading, Manipulating and Writing Data

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Outline for this Training

- Introduction to R and R Studio
- Data Types and Operators
 - Case Study on R Scripting
- Reading, Manipulating and Writing Data
 - Case Study on Dataset Analysis with ETL
- Basic R Programming
 - Case Study: Writing Functions
- Graphics and Plotting
- Deploying R and Dashboard Generation
 - Case Study: Deploying a Simple Dashboard
- Deploying R with C#
 - Case Study: A Simple Standalone GUI For R Apps



Outline for this Session

- Data Frames
- Database Queries
- Reshape Package
- DPLYR Packager



Definition 3.1: Data Frames

- A data frame is a two dimensional dataset.
- Tightly coupled collections of variables which share many of the properties of matrices.
- Used as the fundamental data structure by most of R's modeling scripts



Attributes/Columns/Variables/Features (p + 1)

Tid Refund **Marital Taxable** Cheat **Status** Income Single No Yes 125K 2 Married 100K No No 3 No Single 70K No 4 Yes Married 120K No 5 Divorced Yes No 95K 6 No Married 60K No 220K No 7 Yes Divorced 8 No Single 85K Yes No Married 75K No 9 10 No 90K Yes Single

Rows/ Instances /Tuples /Objects (n)



- Calculating Memory Requirements
 - Given a dataset with 1,500,000 rows and 120 columns, all of which are numeric data. Roughly, how much memory is required to store this data frame?
 - 1,500, 000 rows \times 120 columns \times 8 bytes/numeric entity = 1440000000 bytes
 - = 1440000000 bytes/220 bytes/MB
 - = 1,373.29 MB
 - $= 1.34 \, GB$



- Restrictions on the contents of a data frame
 - The components must be vectors (numeric, character, or logical), factors, numeric matrices, lists, or other data frames.
 - Numeric vectors, logicals and factors are included as is, and by default character vectors are coerced to be factors, whose levels are the unique values appearing in the vector.
 - Vector structures appearing as variables of the data frame must all have the same length, and matrix structures must all have the same row size.



Example 3.1: Data Frames

- > #dataframes
- \triangleright x <- c(10.4, 5.6, 3.1, 6.4, 21.7)
- \triangleright y <- c(11.4, 6.6, 4.1, 7.4, 22.7)
- \triangleright z <- data.frame(x,y)
- \triangleright Z



- To refer to a specific column use the \$ notation.
- Cumbersome
- Use attach() and detach() to make the components of a data frame temporarily visible as variables under their component name



Example 3.2: Attach and Detach

- #dataframes using \$ symbol
- \rightarrow a=z\$x+z\$y
- > a
- > #dataframes using attach and detach
- > attach(z)
- \triangleright b=x+y
- > b
- > detach(z)



```
> #dataframes using $ symbol
> a=z$x+z$y
> a
[1] 21.8 12.2 7.2 13.8 44.4
> #dataframes using attach and detach
> attach(z)
The following objects are masked _by_ .GlobalEnv:
   х, у
> b=x+y
> b
[1] 21.8 12.2 7.2 13.8 44.4
> detach(z)
>
```



- Reading Data from Files
 - Large data objects will usually be read as values from external files rather than entered during an R session at the keyboard.
 - R input facilities are simple and their requirements are fairly strict and even rather inflexible.
 - Types
 - Read from CSV/Text Files
 - Read from Excel Files
 - Read from Database Files



Read from CSV Files

- The sample data can also be in comma separated values (CSV) format.
- Each cell inside such data file is separated by a special character, which usually is a comma, although other characters can be used as well.
- Reads files from the current working directory
- read.csv(file="filename.csv", head=TRUE, sep=",")



- Copy Data from the Desktop
 - For the following exercises, please copy the contents from the
 Shared Folder given to you to the My Documents -> Work folder
 - Data files to be read need to be in the current working directory for ease of access
 - Otherwise, the directory needs to be explicitly stated.
 - read.csv(file="C:/My
 Documents/filename.csv", head=TRUE, sep=",")



Example 3.3: Reading a CSV File

- > #read a CSV file
- heisenberg <read.csv(file="simple.csv",head=TRUE,se
 p=",")</pre>
- > heisenberg



Read from Excel Files

- We can use the function loadWorkbook from the XLConnect package to read the entire workbook
- Then load the worksheets with readWorksheet.
- However, the XLConnect package requires Java to be pre-installed.



Example 3.4: Reading an Excel File

- > #Read from Excel
- ▶ library(xlsx)
- heisenbergxls = read.xlsx("simple.xlsx",
 sheetName = "simple")



- Reading Larger Files
 - An alternative to reading large CSV files
 - Similar to read.csv but faster
 - Use the data.table package



Example 3.5: Read from a large file

- > #Read Large CSV File
- > library("data.table")
- heisenberglargecsv =
 as.data.frame(fread("largecsv.csv"))

```
> #Read Large CSV File
```

- > library("data.table")
- > heisenberglargecsv = as.data.frame(fread("largecsv.csv"))



Definition 3.2: Subsetting

- R has powerful indexing features for accessing object elements.
- These features can be used to select and exclude variables and observations.
- We define here ways to keep or delete variables and observations and to take random samples from a dataset.
- Syntax
 Dataset[select rows , select columns]



Example 3.6: Subsetting

- #Subsetting
- heisenbergaonly =
 heisenberg[heisenberg\$trial=="A",c("tri
 al","mass")]
- > heisenbergaonly

Definition 3.3: Merging

- To merge two data frames (datasets) horizontally, use the merge function.
- In most cases, you join two data frames by one or more common key variables (i.e., an inner join).
 - Inner join: merge(x = df1, y = df2, by = "colname")
 - Outer join: merge(x = df1, y = df2, by = "colname", all = TRUE)
 - Left outer: merge(x = df1, y = df2, by = "colname", all.x = TRUE)
 - Right outer: merge(x = df1, y = df2, by = "colname", all.y = TRUE)



Example 3.7: Merging

- > trial <- c("A", "C", "D")
- \triangleright cost <- c(11.4, 3.3, 1.1)
- trialcost <- data.frame(trial,cost)</pre>
- > trialcost



Example 3.7 (Cont.): Merging

- > #merge
- innerjoin =
 merge(x=heisenberg,y=trialcost,by=c("trial"))
- outerjoin =
 merge(x=heisenberg,y=trialcost,by=c("trial"), all=
 TRUE)
- P leftjoin =
 merge(x=heisenberg,y=trialcost,by=c("trial"),
 all.x=TRUE)
- rightjoin =
 merge(x=heisenberg,y=trialcost,by=c("trial"),
 all.y=TRUE)



heisenberg

trial	mass	velocity
Α	10.0	12
Α	11.0	14
В	5.0	8
В	6.0	10
Α	10.5	13
В	7.0	11

merge

trialcost

trial	cost
Α	11.4
C	3.3
D	1.1

Inner Join

trial	mass	velocity	cost
Α	10.0	12	11.4
Α	11.0	14	11.4
Α	10.5	13	11.4

Outer Join

trial	mass	velocity	cost
Α	10.0	12	11.4
Α	11.0	14	11.4
Α	10.5	13	11.4
В	5.0	8	NA
В	6.0	10	NA
В	7.0	11	NA
С	NA	NA	3.3
D	NA	NA	1.1

Left Join

trial	mass	velocity	cost
Α	10.0	12	11.4
Α	11.0	14	11.4
Α	10.5	13	11.4
В	5.0	8	NA
В	6.0	10	NA
В	7.0	11	NA

Right Join

trial	mass	velocity	cost
Α	10.0	12	11.4
Α	11.0	14	11.4
Α	10.5	13	11.4
С	NA	NA	3.3
D	NA	NA	1.1

- To sort a data frame in R, use the order() function.
- By default, sorting is ASCENDING. Prepend the sorting variable by a minus sign to indicate DESCENDING order.



Example 3.8: Sorting

- > #Sort by Mass
- heisenbergmass = heisenberg[order(heisenberg\$mass),]
- heisenbergmass
- > #Sort by Trial then By Mass
- heisenbergtrialmass=
 heisenberg[order(heisenberg\$trial,
 heisenberg\$mass),]
- heisenbergtrialmass
- > #Sort by Trial then by Mass Descending
- heisenbergtrialmassdesc= heisenberg[order(
 heisenberg\$trial, -heisenberg\$mass),]
 - heisenbergtrialmassdesc



```
> #Sort by Mass
> heisenbergmass = heisenberg[order(heisenberg$mass),]
> heisenbergmass
 trial mass velocity
3
     B 5.0
4
     B 6.0
6
     B 7.0
                  11
1
     A 10.0
              12
5
     A 10.5
            13
     A 11.0
                  14
> #Sort by Trial then By Mass
 heisenbergtrialmass= heisenberg[order(heisenberg$trial, heisenberg$mass),]
> heisenbergtrialmass
 trial mass velocity
     A 10.0
     A 10.5
                  13
     A 11.0
                  14
       5.0
       6.0
                  10
       7.0
                  11
```

Outline for this Session

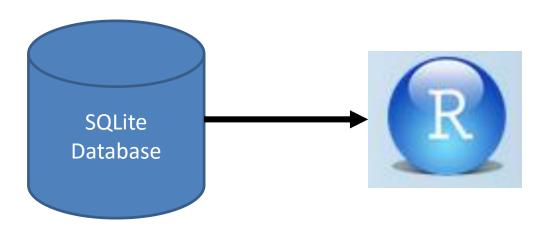
- Data Frames
- Database Queries
- Reshape Package
- DPLYR Packager



- Reading from an SQLite Database
- Reading from an MySQL Database



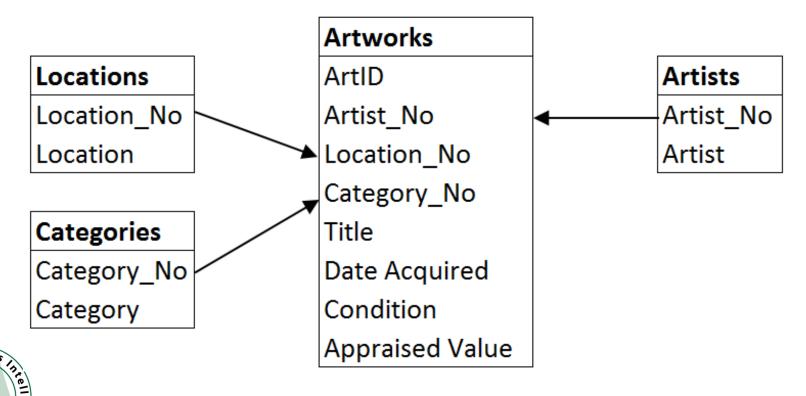
- Reading from an SQLite Database
 - SQLite is an open source database file
 - Connecting to an SQLite Database file is made very easy with the sqlLdf package





Example 3.9: Art Database

SQLite Database Schema: Art Database



- library(sqldf)
- library(XLConnect)
- b db <- dbConnect(SQLite(),
 dbname="Art.sqlite")</pre>
- > rs = dbSendQuery(db, "SELECT * FROM artworks")
- \triangleright data = fetch(rs, n=-1)
- dbDisconnect(db)
- > data



```
> library(sqldf)
> library(XLConnect)
> db <- dbConnect(SQLite(), dbname="Art.sqlite")</pre>
> rs = dbSendQuery(db, "SELECT * FROM artworks")
> data = fetch(rs, n=-1)
> dbDisconnect(db)
[1] TRUE
Warning message:
In .local(conn, ...) : Closing open result set
> data
   ArtID Artist_No Category_No Location_No
                                                                     Title Date.Acquired
                                                        Red Rock Mountain
                                                                                 3/19/05
                                                               Offerings
                                                                                 5/16/05
3
                                                           Spring Flowers
                                                                                 3/20/04
                                                          Seeking Shelter
                                                                                 10/8/05
4
                                                                 The Hang
                                                                                 7/16/04
                                                         House Remembered
                                                                                 8/16/04
6
```



- Reading from an MySQL Database
 - Connecting to MySQL is made very easy with the RMySQL package
 - Once the RMySQL library is installed create a database connection object.
 - mydb = dbConnect(MySQL(), user='user',
 password='password', dbname='database_name',
 host='host')
 - To retrieve data from the database we need to save a results set object.
 - rs = dbSendQuery(mydb, "select * from some table")



Exporting Data

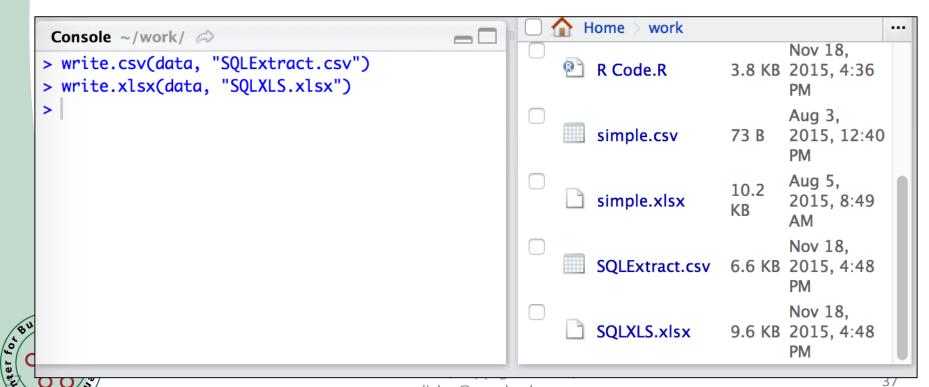
- There are numerous methods for exporting R objects into other formats.
- write.csv(datavariable, "filename.csv")
- write.xlsx(datavariable, "filename.xlsx")



Database Queries

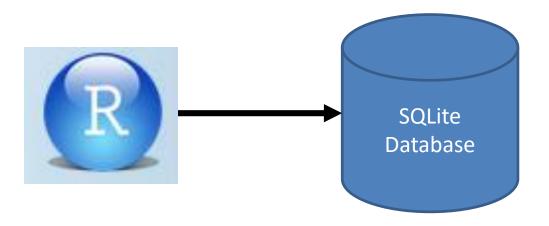
Example 3.10: Exporting Data

- write.csv(data, "SQLExtract.csv")
- write.xlsx(data, "SQLXLS.xlsx")



Database Queries

- Exporting Data to MySQL
 - Write a local data frame or file to the database.





Database Queries

Example 3.11: Exporting to an SQLite Database

- b db <- dbConnect(SQLite(),
 dbname="Art.sqlite")</pre>
- b dbWriteTable(conn = db, name =
 "artworks2", value =data, row.names =
 FALSE, append = TRUE)
- > rs = dbSendQuery(db, "SELECT * FROM artworks2")
- \triangleright datatest = fetch(rs, n=-1)
- dbDisconnect(db)



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Definition 3.2: Reshape

- Reshaping data is a common task in practical data analysis, and it is usually tedious and unintuitive.
- Data often has multiple levels of grouping (nested treatments, split plot designs, or repeated measurements) and typically requires investigation at multiple levels.
- Data reshaping is easiest to define with respect to aggregation.
- Aggregation is a common and familiar task where data is reduced and rearranged into a smaller, more convenient form, with a concomitant reduction in the amount of information

Definition 3.3: Melt

 Melting is the process of transforming measures/facts into a single column/variable

Melt by Mass and Velocity Example

trial mass velocity 10.0 Α 12 14 Α 11.0 dimensions 5.0 8 6.0 В 10 10.5 13 В 7.0 11

facts

Melt

trial variable value 10.0 Α mass 11.0 mass В 5.0 mass 6.0 mass 10.5 Α mass 7.0 mass velocity 12.0 Α velocity 14.0 velocity 8.0 В В velocity 10.0 velocity Α 13.0 В velocity 11.0



Definition 3.4: Cast

- Casting is the summarization of a molten dataset into aggregated data
- Get mean of mass and velocity by trial

trial	variable	value			
Α	mass	10.0			
Α	mass	11.0			
В	mass	5.0	0 1		
В	mass	6.0	Cast	trial	mass
Α	mass	10.5		Α	10.5
В	mass	7.0		В	6.0
Α	velocity	12.0			
Α	velocity	14.0			
В	velocity	8.0			
В	velocity	10.0			
Α	velocity	13.0	E.D. L. Jaloo Conveight IID NEC		
R	velocity	11 0	E.R. L. Jalao, Copyright UP NEC	,	



velocity

13.000000

9.666667

Example 3.12: Reshaping

- library("reshape2")
- heisenberg.m = melt(heisenberg, id=c('trial'), measure=c('mass','velocity'))
- heisenberg.m
- heisenberg.c = dcast(heisenberg.m, trial ~ variable, mean)
- heisenberg.c



```
> library("reshape2")
> heisenberg.m = melt(heisenberg, id=c('trial'), measure=c('mass','velocity'))
> heisenberg.m
  trial variable value
      Α
            mass 10.0
1
      Α
            mass 11.0
3
                   5.0
            mass
                   6.0
            mass
            mass 10.5
            mass 7.0
      A velocity 12.0
      A velocity 14.0
9
      B velocity 8.0
10
      B velocity 10.0
      A velocity 13.0
11
12
       B velocity 11.0
> heisenberg.c = dcast(heisenberg.m, trial ~ variable, mean)
> heisenberg.c
  trial mass velocity
     A 10.5 13.000000
     B 6.0 9.666667
>
```



Outline for this Session

- Data Frames
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Definition 3.5: dplyr Package

- dplyr is a new package which provides a set of tools for efficiently manipulating datasets in R.
- dplyr is the next iteration of plyr, focussing on only data frames.
- dplyr is faster, has a more consistent API and should be easier to use.
- Succeeding examples taken from: <u>https://cran.rstudio.com/web/packages/dplyr/vignettes/introduction.html</u>



- The nycflights13 data frame.
 - This dataset contains all 336776 flights that departed from and arrived at New York City in 2013. The data comes from the US Bureau of Transportation Statistics, and is documented in ?nycflights13



- The filter() function
 - Allows you to select a subset of rows in a data frame. The first argument is the name of the data frame.
 - The second and subsequent arguments are the expressions that filter the data frame



Example 3.13a: Filtering

- Select all flights last January 1, 2013
- library("dplyr")
- flights = read.csv("flights.csv")
- filtered = filter(flights, month == 1, day == 1)
- View(filtered)



Example 3.13 (Cont.): Filtering

	Х	year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	hot ^
1	1	2013	1	1	517	515	2	830	819	11	UA	1545	N14228	EWR	IAH	227	1400	5
2	2	2013	1	1	533	529	4	850	830	20	UA	1714	N24211	LGA	IAH	227	1416	5
3	3	2013	1	1	542	540	2	923	850	33	AA	1141	N619AA	JFK	MIA	160	1089	5
4	4	2013	1	1	544	545	-1	1004	1022	-18	B6	725	N804JB	JFK	BQN	183	1576	5
5	5	2013	1	1	554	600	-6	812	837	-25	DL	461	N668DN	LGA	ATL	116	762	6
6	6	2013	1	1	554	558	-4	740	728	12	UA	1696	N39463	EWR	ORD	150	719	5
7	7	2013	1	1	555	600	-5	913	854	19	B6	507	N516JB	EWR	FLL	158	1065	6
8	8	2013	1	1	557	600	-3	709	723	-14	EV	5708	N829AS	LGA	IAD	53	229	6
9	9	2013	1	1	557	600	-3	838	846	-8	B6	79	N593JB	JFK	MCO	140	944	6
10	10	2013	1	1	558	600	-2	753	745	8	AA	301	N3ALAA	LGA	ORD	138	733	6
11	11	2013	1	1	558	600	-2	849	851	-2	B6	49	N793JB	JFK	PBI	149	1028	6
12	12	2013	1	1	558	600	-2	853	856	-3	B6	71	N657JB	JFK	TPA	158	1005	6
13	13	2013	1	1	558	600	-2	924	917	7	UA	194	N29129	JFK	LAX	345	2475	6
14	14	2013	1	1	558	600	-2	923	937	-14	UA	1124	N53441	EWR	SF0	361	2565	6
15	15	2013	1	1	559	600	-1	941	910	31	AA	707	N3DUAA	LGA	DFW	257	1389	6
16	16	2013	1	1	559	559	0	702	706	-4	B6	1806	N708JB	JFK	BOS	44	187	5



Example 3.13b: Filtering

- Select all flights from January or July 2013.
- \triangleright filtered2 = filter(flights, month == 1 | month == 7)
- View(filtered2)



Example 3.13 (Cont.): Filtering

	X	year [‡]	month ‡	day [‡]	dep_time ‡	sched_dep_time ‡	dep_delay [‡]	arr_time ‡	sched_arr_time ‡	arr_delay [‡]	carrier	÷
56420	279866	2013	7	31	2243	2245	-2	2348	1	-13	B6	^
56421	279867	2013	7	31	2245	2245	0	2400	4	-4	B6	
56422	279868	2013	7	31	2248	2250	-2	5	14	-9	B6	
56423	279869	2013	7	31	2325	2051	154	221	2358	143	B6	
56424	279870	2013	7	31	2346	2305	41	38	13	25	B6	
56425	279871	2013	7	31	2352	2245	67	49	2359	50	B6	
56426	279872	2013	7	31	NA	655	NA	NA	930	NA	AA	
56427	279873	2013	7	31	NA	1400	NA	NA	1508	NA	US	
56428	279874	2013	7	31	NA	959	NA	NA	1125	NA	UA	
56429	279875	2013	7	31	NA	1025	NA	NA	1225	NA	MQ	~



- The slice () function
 - To select rows by position, use slice():
 - slice(df, rowindeces)



Example 3.14: Slice

- Select first 10 rows of the flights dataset.
- > slice(flights, 1:10)

	X ‡	year [‡]	month ‡	day [‡]	dep_time [‡]	sched_dep_time +	dep_delay [‡]	arr_time ‡	sched_arr_time ‡
1	1	2013	1	1	517	515	2	830	819
2	2	2013	1	1	533	529	4	850	830
3	3	2013	1	1	542	540	2	923	850
4	4	2013	1	1	544	545	-1	1004	1022
5	5	2013	1	1	554	600	-6	812	837
6	6	2013	1	1	554	558	-4	740	728
7	7	2013	1	1	555	600	-5	913	854
8	8	2013	1	1	557	600	-3	709	723
9	9	2013	1	1	557	600	-3	838	846
10	10	2013	1	1	558	600	-2	753	745

- Use the arrange () function to sort rows
 - arrange () works similarly to filter () except that instead of filtering or selecting rows, it reorders them.
 - It takes a data frame, and a set of column names (or more complicated expressions) to order by. If you provide more than one column name, each additional column will be used to break ties in the values of preceding columns:



Example 3.15: Arrange

- Sort by year, then by month and then by day.
- sorted = arrange(flights, year, month, day)
- View(sorted)

	Х	year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	h
1	1	2013	1	1	517	515	2	830	819	11	UA	1545	N14228	EWR	IAH	227	1400	5
2	2	2013	1	1	533	529	4	850	830	20	UA	1714	N24211	LGA	IAH	227	1416	5
3	3	2013	1	1	542	540	2	923	850	33	AA	1141	N619AA	JFK	MIA	160	1089	5
4	4	2013	1	1	544	545	-1	1004	1022	-18	B6	725	N804JB	JFK	BQN	183	1576	5
5	5	2013	1	1	554	600	-6	812	837	-25	DL	461	N668DN	LGA	ATL	116	762	6
6	6	2013	1	1	554	558	-4	740	728	12	UA	1696	N39463	EWR	ORD	150	719	5
7	7	2013	1	1	555	600	-5	913	854	19	B6	507	N516JB	EWR	FLL	158	1065	6
8	8	2013	1	1	557	600	-3	709	723	-14	EV	5708	N829AS	LGA	IAD	53	229	6
9	9	2013	1	1	557	600	-3	838	846	-8	B6	79	N593JB	JFK	MCO	140	944	6
10	10	2013	1	1	558	600	-2	753	745	8	AA	301	N3ALAA	LGA	ORD	138	733	6



Example 3.15 (Cont.): Arrange

- Sort by descending arrival delay time
- descsorted = arrange(flights, desc(arr delay))
- View(descsorted)

$\langle \! \downarrow \! \rangle$	336,776 observations of 20 variables																	
	X	year	month	day	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum	origin	dest	air_time	distance	^
1	7073	2013	1	9	641	900	1301	1242	1530	1272	HA	51	N384HA	JFK	HNL	640	4983	
2	235779	2013	6	15	1432	1935	1137	1607	2120	1127	MQ	3535	N504MQ	JFK	CMH	74	483	
3	8240	2013	1	10	1121	1635	1126	1239	1810	1109	MQ	3695	N517MQ	EWR	ORD	111	719	
4	327044	2013	9	20	1139	1845	1014	1457	2210	1007	AA	177	N338AA	JFK	SF0	354	2586	
5	270377	2013	7	22	845	1600	1005	1044	1815	989	MQ	3075	N665MQ	JFK	CVG	96	589	
6	173993	2013	4	10	1100	1900	960	1342	2211	931	DI	2391	NGSGDI	JEK	ТРΔ	139	1005	



- The select () Function to select certain columns
 - Selecting Columns from Datasets by Listing Column Names
 - select(df, col1, col2, col3,...)
 - Selecting all columns between two columns (inclusive)
 - select(df, col2:col5)
 - Excluding all columns between two columns (inclusive) and selecting all others
 - select(df, -col2:col5)



Example 3.16: Select

- Select only Year, Month and Day columns
- > selectedcol = select(flights, year, month, day)
- View(selectedcol)

	year	month	day
1	2013	1	1
2	2013	1	1
3	2013	1	1
4	2013	1	1
5	2013	1	1
6	2013	1	1



Example 3.16 (Cont.): Select

- Select all rows except columns on Year, Month and Day
- > selectedcol = select(flights, -(year:day))
- > View(selectedol)

	X	dep_time	sched_dep_time	dep_delay	arr_time	sched_arr_time	arr_delay	carrier	flight	tailnum
1	1	517	515	2	830	819	11	UA	1545	N14228
2	2	533	529	4	850	830	20	UA	1714	N24211
3	3	542	540	2	923	850	33	AA	1141	N619AA



- Extract Distinct (unique) Rows
 - Use distinct() to find unique values in a table



Example 3.17: Distinct

- Select unique Origins and Destinations
- distinct = distinct(select(flights, origin, dest))
- > View(distinct)

origin	dest
EWR	IAH
LGA	IAH
JFK	MIA
JFK	BQN
LGA	ATL
EWR	ORD
EWR	FLL
LGA	IAD
JFK	MCO
LGA	ORD
	EWR LGA JFK LGA EWR EWR LGA JFK



- Adding New Columns With mutate ()
 - Besides selecting sets of existing columns, it's often useful to add new columns that are functions of existing columns. This is the job of mutate():
 - Mutate allows you to refer to columns that was just created
 - mutate(df, newcol1 = f(oldcols), newcol2 = f(newcol1))



Example 3.18: Mutate

Calculate new columns on gain, speed and gain per hour.

			Q,					
\$	gain ‡	speed ‡	gain_per_hour ‡					
00	9	370.0441	2.3788546					
0	16	374.2731	4.2290749					
0	31	408.3750	11.6250000					
0	-17	516.7213	-5.5737705					
0	-19	394.1379	-9.8275862					
0	16	287.6000	6.4000000					
00	24	404.4304	9.1139241					



- Summarize values with summarise()
 - It collapses a data frame to a single row
 - summarise(df, var = mean(cols))



Example 3.19: Summarize

- Calculate mean delay
- meandelay = summarise(flights,
- delay = mean(dep_delay, na.rm = TRUE))
- View (meandelay)

	delay [‡]
1	12.63907



- dplyr verbs are useful on their own, but they become really powerful when you apply them to groups of observations within a dataset.
- In dplyr, this is done by with the group by () function.
- It breaks down a dataset into specified groups of rows.
- When the verbs are applied on the resulting object they'll be automatically applied "by group". Most importantly, all this is achieved by using the same exact syntax you'd use with an ungrouped object.



- grouped select() is the same as ungrouped select(),
 except that grouping variables are always retained.
- grouped arrange() orders first by the grouping variables
- mutate() and filter() are most useful in conjunction with window functions (like rank(), or min(x) == x).
- slice() extracts rows within each group.
- summarise() can mimic the reshape function as shows in the succeeding examples



Example 3.20: Grouping

- For each unique plane tail number, count the number of flights it made, average distance travelled and average delay time.
- by tailnum <- group by(flights, tailnum)</pre>
- delay <- summarise(by tailnum,</pre>
- \triangleright count = n(),
- dist = mean(distance, na.rm = TRUE),
- delay = mean(arr_delay, na.rm = TRUE))
- delay <- filter(delay, count > 20, dist < 2000)</pre>
- View(delay)



Example 3.20 (Cont.): Grouping

	tailnum [‡]	count ‡	dist [‡]	delay [‡]
1	N0EGMQ	371	676.1887	9.9829545
2	N10156	153	757.9477	12.7172414
3	N102UW	48	535.8750	2.9375000
4	N103US	46	535.1957	-6.9347826
5	N104UW	47	535.2553	1.8043478
6	N10575	289	519.7024	20.6914498
7	N105UW	45	524.8444	-0.2666667

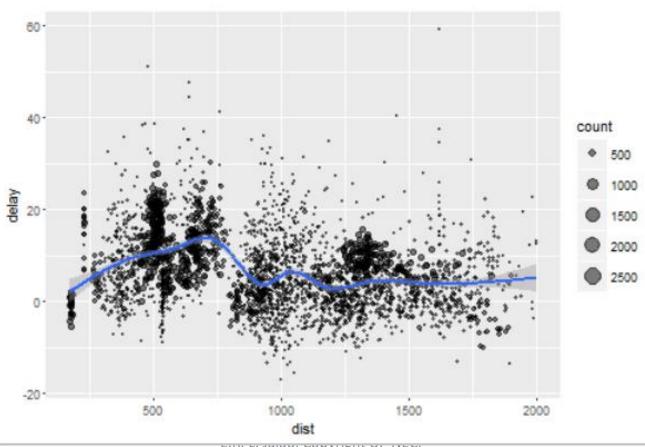


Example 3.21: Grouping with Visualization

- Plot Distance Versus Delay
- library("ggplot2")
- ggplot(delay, aes(dist, delay)) +
- \triangleright geom point(aes(size = count), alpha = 1/2) +
- peom_smooth() +
- scale size area()



Example 3.21 (Cont.): Grouping with Visualization



- Summarize() and aggregate functions can be used together, which take a vector of values and return a single number.
- There are many useful examples of such functions in base R like min(), max(), mean(), sum(), sd(), median(), and IQR().
- dplyr provides a handful of others:
 - -n (): the number of observations in the current group
 - n distinct (x): the number of unique values in x.
 - first(x), last(x) and nth(x, n)



Example 3.22: Summarize and Aggregate

- For each destination, calculate number of planes, flights and average delay
- destinations <- group by(flights, dest)</pre>
- destsummary = summarise(destinations,

E.R.

- \triangleright planes = n distinct(tailnum),
- \triangleright flights = n(),
- delay = mean(dep_delay, na.rm = TRUE))
- View(destsummary)

	dest ‡	planes ‡	flights ‡	delay [‡]
1	ABQ	108	254	13.740157
2	ACK	58	265	6.456604
3	ALB	172	439	23.620525
4	ANC	6	8	12.875000
5	ATL	1180	17215	12.509824
6	AUS	993	2439	13.025641



Example 3.22 (Cont.): Summarize and Aggregate

	dest ‡	planes [‡]	flights ‡	delay [‡]
1	ABQ	108	254	13.740157
2	ACK	58	265	6.456604
3	ALB	172	439	23.620525
4	ANC	6	8	12.875000
5	ATL	1180	17215	12.509824
6	AUS	993	2439	13.025641



Chaining

- The dplyr API is functional in the sense that function calls don't have side-effects.
- Results must be saved often. This doesn't lead to particularly elegant code, especially if many operations are needed.
- dplyr provides the %>% operator.
 - $x \approx >$ f(y) turns into f(x, y) so you can use it to rewrite multiple operations that you can read left-to-right, top-to-bottom



Example 3.23: Chaining

```
delayreport=flights %>%
group by (year, month, day) %>%
    select(arr delay, dep delay) %>%
summarise(
arr = mean(arr delay, na.rm = TRUE),
dep = mean(dep delay, na.rm = TRUE)
filter(arr > 30 \mid dep > 30)
```



<pre>View(delayreport)</pre>		year ‡	month ‡	day [‡]	arr ‡	dep [‡]
	1	2013	1	16	34.24736	24.61287
100	2	2013	1	31	32.60285	28.65836
	3	2013	2	11	36.29009	39.07360
E.R. L.	4	2013	2	27	31.25249	37.76327

Case 2

- Case Study on Dataset Analysis with ETL
- You are to build a simple ETL process for the Art Database



References

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