SAS <-> R :: CHEAT SHEET

Introduction

This guide aims to familiarise SAS users with R.
R examples make use of tidyverse collection of packages.
Install tidyverse:
Attach tidyverse packages for use:

Install.packages("tidyverse")
Iibrary(tidyverse)

R data here in 'data frames', and occasionally vectors (via **c()**) Other R structures (lists, matrices...) are not explored here.

Keyboard shortcuts: <- Alt + - %>% Ctrl + Shift + m

Datasets; drop, keep & rename variables

```
data new data:
                                  new data <- old data
 set old data;
run;
data new data (keep=id) ;
                                  new data <- old data %>%
set old data (drop=job title);
                                   select(-job title) %>%
                                   select(id)
data new data (drop = temp: );
                                  new data <- old data %>%
set old_data;
                                   select( -starts with("temp") )
run:
                                              C.f. contains(), ends with()
                                  new data <- old data %>%
data new data:
 set old data:
                                   rename(new_name = old_name)
 rename old name = new name;
                                                       Note order differs
```

Conditional filtering

```
data new data;
                                    new data <- old data %>%
 set old data;
                                     filter(Sex == "M")
 if Sex = "M":
run:
data new data:
                                    new data <- old data %>%
                                     filter(year %in% c(2010,2011,2012))
set old data:
 if year in (2010,2011,2012);
run;
data new data;
                                    new data <- old data %>%
 set old data;
                                     group_by( id ) %>%
 bv id:
                                     slice(1)
 if first.id:
run:
                                                 Could use slice(n( )) for last
                                    new data <- old data %>%
data new data:
                                     filter(dob > as.Date("1990-04-25"))
 set old data:
 if dob > "25APR1990"d;
run;
```

New variables, conditional editing

```
data new data;
                                        new data <- old data %>%
                                         mutate(total income = wages + benefits)
 set old data;
 total_income = wages + benefits;
data new_data;
                                        new_data <- old_data %>%
 set old data:
                                         mutate(full time = if else(hours > 30."Y", "N"))
 if hours > 30 then full_time = "Y";
 else full time = "N";
data new data;
                                        new data <- old data %>%
 set old data;
                                         mutate(weather = case when(
 if temp > 20 then weather = "Warm":
                                           temp > 20 ~ "Warm".
                                          temp > 10 ~ "Mild",
 else if temp > 10 then weather = "Mild";
 else weather = "Cold":
                                           TRUE ~ "Cold" ))
```

Counting and Summarising

```
proc freq data=old data;
                                         old data %>%
 table job_type;
                                          count( job_type )
                                                                           For percent, add:
                                                         %>% mutate(percent = n*100/sum(n))
proc freq data=old data;
                                         old data %>%
                                          count( job_type , region )
 table job_type*region;
proc summary data=old_data nway;
                                         new_data <- old_data %>%
                                          group_by(( job_type , region ) %>%
 class job_type region;
 output out = new data;
                                          summarise( Count = n())
                                                 Equivalent without nway not trivially produced
proc summary data=old data nway;
                                         new data <- old data %>%
 class job type region;
                                          group by(job type, region) %>%
 var income;
                                          summarise( total_salaries = sum( salary ) ,
 output out = new data
                                             Count = n()
   sum( salary ) = total_salaries;
                                                  Lots of summary functions in both languages
                            Swap summarise() for mutate() to add summary data to original data
```

Combining datasets

```
data new_data; new_data <- bind_rows( data_1 , data_2 )

set data_1 data_2 ;
run;

data new_data; new_data <- full_join( data_1 , data_2 , by = "id")

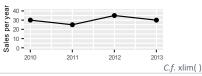
merge data_1 data_2 ;
by id ;
run; C.f. left_join() , right_join() , inner_join()
```

Some plotting in R

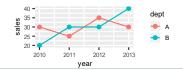
```
ggplot( my_data , aes( year , sales ) ) +
geom_point( ) + geom_line( )

35.0 -
g 32.5 -
g 30.0 -
27.5 -
25.0 -
2010 2011 2012 2013
```

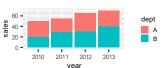
```
ggplot( my_data , aes( year , sales ) ) +
geom_point( ) + geom_line( ) + ylim(0, 40)
labs(x="" , y="Sales per year")
```



ggplot(my_data , aes(year , sales, **colour=**dept)) + geom point() + geom line()



ggplot(my_data , aes(year , sales, fill=dept)) +
geom_col()



N.B. 'colour' for lines & points, 'fill' for shapes

```
ggplot( my_data , aes( year , sales, fill=dept ) ) +
geom_col( position="dodge" ) + coord_flip( )
```



C.f. position = "fill" for 100% stacked bars/cols

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Sorting and Row-Wise Operations

```
proc sort data=old data out=new data:
                                              new data <- old data %>%
 by id descending income;
                                               arrange( id , -income )
run;
proc sort data=old_data nodup;
                                              old data <- old data %>%
by id job_type;
                                               arrange(id, job_type)%>%
run;
                                               distinct()
                                  N.B. nodup relies on adjacency of duplicate rows, distinct() does not
proc sort data=old_data nodupkey;
                                              old data <- old data %>%
by id;
                                               arrange(id)%>%
run:
                                               group_by( id ) %>%
                                               slice(1)
data new data:
                                              new_data <- old_data %>%
 set old data;
                                               group by( id ) %>%
 by id descending income;
                                             slice(which.max( income ))
                                                                                C.f. which.min()
 if first.id;
                                 Swap to preserve duplicate maxima: ... filter(income == max(income))
run:
                                                               alternatively: ... top n(1, income)
data new data;
                                              new data <- old data %>%
 set old data:
                                               mutate( prev_id= lag( id, 1 ))
 prev_id= lag( id );
                                                                   C.f. lead() for subsequent rows
data new data:
                                              new_data <- old_data %>%
 set old data;
                                               group by( id ) %>%
                                               mutate( counter = row_number( ) )
 by id;
 counter + 1;
 if first.id then counter = 1:
```

Converting and Rounding

```
data new_data;
                                             new data <- old data %>%
 set old data;
                                              mutate(num_var = as.numeric( "5" )) %>%
 num var= input("5", 8.);
                                              mutate(text var = as.character( 5 ))
 text_var = put(5 , 8.);
run;
data new data :
                                             new data <- old data %>%
 set old data:
                                              mutate(nearest 5 = round(x / 5)*5) %>%
 nearest_5 = round(x, 5)
                                              mutate(two_decimals = round( x , digits = 2)
 two decimals = round(x, 0.01)
run;
```

Creating functions to modify datasets

Dealing with strings

```
data new_data;
                                                      new data <- old data %>%
                                                       filter(str detect( job title , "Health"))
 set old data;
if find( job_title , "Health");
data new data;
                                                      new data <- old data %>%
                                                       filter(str_detect( job_title , "^Health"))
set old data;
 if job_title =: "Health";
run;
                                                                         Use ^ for start of string, $ for end of string, e.g. "Health$"
data new data:
                                                      new_data <- old_data %>%
set old data:
                                                       mutate( substring = str_sub( big_string , 3 , 6 ))
 substring = substr( big string , 3, 4);
                                                    Returns characters 3 to 6. Note SAS uses <start> <length>, R uses <start>, <end>
data new data;
                                                      new data <- old data %>%
set old data;
                                                       mutate( address = str_replace_all( address , "Street", "St" ))
 address = tranwrd( address, "Street", "St");
run:
                                                                                C.f. str_replace() for first instance of pattern only
                                                      new data <- old data %>%
data new_data;
set old_data;
                                                       mutate( full_name = str_c( first_name , surname , sep =" "))
 full_name = catx(" ", first_name, surname);
                                                                                     Drop sep = " " for equivalent to cats( ) in SAS
                                                      new data <- old data %>%
data new data;
 set old data;
                                                       mutate( first word= word(sentence , 1 ))
 first word= scan( sentence , 1);
                                                                 R example preserves punctuation at the end of words, SAS doesn't
data new data:
                                                      new data <- old data %>%
                                                       mutate( house_number = str_extract( address, "\\d*" ))
set old data:
house_number = compress( address , , "dk" );
run:
                                                         Wide range of regexps in both languages, this example extracts digits only
```

File operations

proc import datafile = "my_file.csv"

out = my data dbms = csv;

```
Operate in 'Work' library.
                                                      Operate in a particular 'working directory' (identify using getwd())
Use libname to define file locations
                                                      Move to other locations using setwd()
libname library name "file location";
                                                      save(data in use , file="file location/saved data.rda")
data library name.saved data:
 set data_in_use;
                                                      setwd("file_location")
run:
                                                      save(data in use . file="saved data.rda")
libname library name "file location":
                                                      load( "file_location/saved_data.rda" )
data data_in_use;
 set library_name.saved_data;
                                                      setwd("file_location")
                                                                                        save() can store multiple data frames in a
                                                      load("saved data.rda")
                                                                                     single .rda file, load() will restore all of these
```

my_data <- read_csv("my_file.csv")

Both examples assume columns headers in csv file