SAS <-> R :: CHEAT SHEET

Introduction

This guide aims to familiarise SAS users with R. R examples make use of tidyverse collection of packages. Install tidvverse: install.packages("tidyverse") library(tidyverse) Attach tidyverse packages for use: 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

```
new data <- old data
data new data;
 set old data;
run:
data new data (keep=id);
                                  new data <- old data %>%
 set old_data (drop=job_title) ;
                                    select(-iob title) %>%
run:
                                    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:
                                    rename(new name = old name)
 set old data;
 rename old name = new name;
                                                        Note order differs
```

Conditional filtering

```
new data <- old data %>%
data new_data;
 set old data:
                                     filter(Sex == "M")
if Sex = "M":
run;
data new_data;
                                   new_data <- old_data %>%
 set old data;
                                    filter(year %in% c(2010,2011,2012))
 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;
 set old data;
                                     filter(dob > as.Date("1990-04-25"))
 if dob > "25APR1990"d;
run:
```

New variables, conditional editing

```
data new data;
                                        new data <- old data %>%
 set old data:
                                         mutate(total income = wages + benefits)
 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".
 else if temp > 10 then weather = "Mild";
                                          temp > 10 ~ "Mild",
 else weather = "Cold";
                                           TRUE ~ "Cold" ) )
```

Counting and Summarising

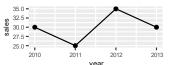
```
proc freq data=old data;
                                         old data %>%
 table job_type;
                                          count( job_type )
                                                                            For percent, add:
run:
                                                         %>% mutate(percent = n*100/sum(n))
proc freq data=old_data;
                                         old data %>%
 table job_type*region;
                                          count( 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
                                          new data <- old data %>%
proc summary data=old data nway;
 class job_type region;
                                          group_by(job_type, region) %>%
                                          summarise( total_salaries = sum( salary ) ,
 var 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

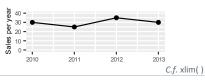
```
data new_data;
                                            new data <- bind rows( data 1, data 2)
 set data_1 data_2;
                                            new_data <- left_join( data_1 , data_2 , by = "id")</pre>
data new data:
 merge data_1 (in = in_1) data_2;
 by id;
 if in 1;
                                                           C.f. full join(), right join(), inner join()
run:
```

Some plotting in R

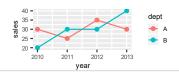
```
ggplot( my_data , aes( year , sales ) ) +
 geom_point() + geom_line()
```



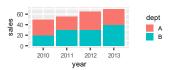
```
ggplot( my_data , aes( year , sales ) ) +
 geom_point() + geom_line() + ylim(0, 40)
 labs(x="", v="Sales per vear")
```



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

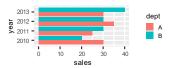


ggplot(my_data , aes(year , sales, fill=dept)) + geom_col()



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

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

```
%macro add variable(dataset name);
                                              add variable <- function ( dataset name ){</pre>
data &dataset name:
                                               dataset name <- dataset name %>%
 set &dataset_name;
                                                mutate(new_variable = 1)
 new_variable = 1;
                                               return( dataset_name )
run;
                                              my_data <- add_variable( my_data )
%mend;
%add_variable( my_data );
                                                            Note SAS can modify within the macro,
                                                       whereas R creates a copy within the function
```

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