R DPLYR Tibble Basics
From Matrixes to Tibble Dataframes

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Go back to fan's REconTools Package, R4Econ Repository, or Intro Stats with R Repository.

```
rm(list = ls(all.names = TRUE))
options(knitr.duplicate.label = 'allow')
```

```
library(tidyverse)
library(knitr)
library(kableExtra)
library(REconTools)
# file name
st_file_name = 'fs_tib_basics'
# Generate R File
purl(paste0(st_file_name, ".Rmd"), output=paste0(st_file_name, ".R"), documentation = 2)
# Generate PDF and HTML
# rmarkdown::render("C:/Users/fan/R4Econ/support/tibble/fs_tib_basics.Rmd", "pdf_document")
# rmarkdown::render("C:/Users/fan/R4Econ/support/tibble/fs_tib_basics.Rmd", "html_document")
```

Tibble Basics

Generate Tibble given Matrixes and Arrays

Given Arrays and Matrixes, Generate Tibble and Name Variables/Columns

- naming tibble columns
- tibble variable names
- dplyr rename tibble
- dplyr rename tibble all variables
- dplyr rename all columns by index
- dplyr tibble add index column
- see also: SO-51205520

```
# Base Inputs
ar_col <- c(-1,+1)
mt_rnorm_a <- matrix(rnorm(4,mean=0,sd=1), nrow=2, ncol=2)
mt_rnorm_b <- matrix(rnorm(4,mean=0,sd=1), nrow=2, ncol=4)

# Combine Matrix
mt_combine <- cbind(ar_col, mt_rnorm_a, mt_rnorm_b)
colnames(mt_combine) <- c('ar_col',
                          paste0('matcolvar_grpa_', seq(1,dim(mt_rnorm_a)[2])),
                          paste0('matcolvar_grpb_', seq(1,dim(mt_rnorm_b)[2])))

# Variable Names
ar_st_varnames <- c('var_one',
```

```r
                    paste0('tibcolvar_ga_', c(1,2)),
                    paste0('tibcolvar_gb_', c(1,2,3,4)))

# Combine to tibble, add name col1, col2, etc.
tb_combine <- as_tibble(mt_combine) %>% rename_all(~c(ar_st_varnames))

# Add an index column to the dataframe, ID column
tb_combine <- tb_combine %>% rowid_to_column(var = "ID")

# Change all gb variable names
tb_combine <- tb_combine %>%
                rename_at(vars(starts_with("tibcolvar_gb_")),
                          funs(str_replace(., "_gb_", "_gbrenamed_")))
```

```
## Warning: funs() is soft deprecated as of dplyr 0.8.0
## Please use a list of either functions or lambdas:
##
##   # Simple named list:
##   list(mean = mean, median = median)
##
##   # Auto named with `tibble::lst()`:
##   tibble::lst(mean, median)
##
##   # Using lambdas
##   list(~ mean(., trim = .2), ~ median(., na.rm = TRUE))
## This warning is displayed once per session.
```

```r
# Tibble back to matrix
mt_tb_combine_back <- data.matrix(tb_combine)

# Display
kable(mt_combine) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

ar__col	matcolvar__grpa__1	matcolvar__grpa__2	matcolvar__grpb__1	matcolvar__grpb__2	matcolvar__grpb__3	matc
-1	0.9191448	0.2542336	0.7344035	-0.7505504	0.7344035	
1	-0.9295410	1.0324020	-1.2482721	1.0079655	-1.2482721	

```r
kable(tb_combine) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

ID	var__one	tibcolvar__ga__1	tibcolvar__ga__2	tibcolvar__gbrenamed__1	tibcolvar__gbrenamed__2	tibcolvar__gbre
1	-1	0.9191448	0.2542336	0.7344035	-0.7505504	
2	1	-0.9295410	1.0324020	-1.2482721	1.0079655	

```r
kable(mt_tb_combine_back) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

ID	var__one	tibcolvar__ga__1	tibcolvar__ga__2	tibcolvar__gbrenamed__1	tibcolvar__gbrenamed__2	tibcolvar__gbre
1	-1	0.9191448	0.2542336	0.7344035	-0.7505504	
2	1	-0.9295410	1.0324020	-1.2482721	1.0079655	

Rename Tibble with Numeric Column Names

After reshaping, often could end up with variable names that are all numeric, intgers for example, how to rename these variables to add a common prefix for example.

```r
# Base Inputs
ar_col <- c(-1,+1)
mt_rnorm_c <- matrix(rnorm(4,mean=0,sd=1), nrow=5, ncol=10)
```

```
## Warning in matrix(rnorm(4, mean = 0, sd = 1), nrow = 5, ncol = 10): data length
## [4] is not a sub-multiple or multiple of the number of rows [5]
```

```r
mt_combine <- cbind(ar_col, mt_rnorm_c)
```

```
## Warning in cbind(ar_col, mt_rnorm_c): number of rows of result is not a multiple
## of vector length (arg 1)
```

```r
# Variable Names
ar_it_cols_ctr <- seq(1, dim(mt_rnorm_c)[2])
ar_st_varnames <- c('var_one', ar_it_cols_ctr)

# Combine to tibble, add name col1, col2, etc.
tb_combine <- as_tibble(mt_combine) %>% rename_all(~c(ar_st_varnames))
```

```
## Warning: `as_tibble.matrix()` requires a matrix with column names or a `.name_repair` argument. Using
## This warning is displayed once per session.
```

```r
# Add an index column to the dataframe, ID column
tb_combine_ori <- tb_combine %>% rowid_to_column(var = "ID")

# Change all gb variable names
tb_combine <- tb_combine_ori %>%
                rename_at(
                  vars(num_range('',ar_it_cols_ctr)),
                  funs(paste0("rho", . , 'var'))
                  )

# Display
kable(tb_combine_ori) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

ID	var_one	1	2	3	4	5	6	7	8
1	-1	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546
2	1	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444
3	-1	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551
4	1	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993
5	-1	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546

```r
kable(tb_combine) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

ID	var_one	rho1var	rho2var	rho3var	rho4var	rho5var	rho6var	rho7var	rho8var
1	-1	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546
2	1	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444
3	-1	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551
4	1	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993
5	-1	1.0846444	0.1743551	-0.7493993	0.5366546	1.0846444	0.1743551	-0.7493993	0.5366546

Tibble Row and Column and Summarize

Show what is in the table: 1, column and row names; 2, contents inside table.

```
tb_iris <- as_tibble(iris)
rownames(tb_iris)
```

```
##   [1] "1"   "2"   "3"   "4"   "5"   "6"   "7"   "8"   "9"   "10"  "11"  "12"
##  [13] "13"  "14"  "15"  "16"  "17"  "18"  "19"  "20"  "21"  "22"  "23"  "24"
##  [25] "25"  "26"  "27"  "28"  "29"  "30"  "31"  "32"  "33"  "34"  "35"  "36"
##  [37] "37"  "38"  "39"  "40"  "41"  "42"  "43"  "44"  "45"  "46"  "47"  "48"
##  [49] "49"  "50"  "51"  "52"  "53"  "54"  "55"  "56"  "57"  "58"  "59"  "60"
##  [61] "61"  "62"  "63"  "64"  "65"  "66"  "67"  "68"  "69"  "70"  "71"  "72"
##  [73] "73"  "74"  "75"  "76"  "77"  "78"  "79"  "80"  "81"  "82"  "83"  "84"
##  [85] "85"  "86"  "87"  "88"  "89"  "90"  "91"  "92"  "93"  "94"  "95"  "96"
##  [97] "97"  "98"  "99"  "100" "101" "102" "103" "104" "105" "106" "107" "108"
## [109] "109" "110" "111" "112" "113" "114" "115" "116" "117" "118" "119" "120"
## [121] "121" "122" "123" "124" "125" "126" "127" "128" "129" "130" "131" "132"
## [133] "133" "134" "135" "136" "137" "138" "139" "140" "141" "142" "143" "144"
## [145] "145" "146" "147" "148" "149" "150"
```

```
colnames(tb_iris)
```

```
## [1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Width"  "Species"
```

```
colnames(tb_iris)
```

```
## [1] "Sepal.Length" "Sepal.Width"  "Petal.Length" "Petal.Width"  "Species"
```

```
summary(tb_iris)
```

```
##   Sepal.Length    Sepal.Width     Petal.Length    Petal.Width
##  Min.   :4.300   Min.   :2.000   Min.   :1.000   Min.   :0.100
##  1st Qu.:5.100   1st Qu.:2.800   1st Qu.:1.600   1st Qu.:0.300
##  Median :5.800   Median :3.000   Median :4.350   Median :1.300
##  Mean   :5.843   Mean   :3.057   Mean   :3.758   Mean   :1.199
##  3rd Qu.:6.400   3rd Qu.:3.300   3rd Qu.:5.100   3rd Qu.:1.800
##  Max.   :7.900   Max.   :4.400   Max.   :6.900   Max.   :2.500
##        Species
##  setosa    :50
##  versicolor:50
##  virginica :50
##
##
##
```

Tibble Sorting

- dplyr arrange desc reverse
- dplyr sort

```r
# Sort in Ascending Order
tb_iris %>% select(Species, Sepal.Length, everything()) %>%
  arrange(Species, Sepal.Length) %>% head(10) %>%
  kable() %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
setosa	4.3	3.0	1.1	0.1
setosa	4.4	2.9	1.4	0.2
setosa	4.4	3.0	1.3	0.2
setosa	4.4	3.2	1.3	0.2
setosa	4.5	2.3	1.3	0.3
setosa	4.6	3.1	1.5	0.2
setosa	4.6	3.4	1.4	0.3
setosa	4.6	3.6	1.0	0.2
setosa	4.6	3.2	1.4	0.2
setosa	4.7	3.2	1.3	0.2

```r
# Sort in Descending Order
tb_iris %>% select(Species, Sepal.Length, everything()) %>%
  arrange(desc(Species), desc(Sepal.Length)) %>% head(10) %>%
  kable() %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

Species	Sepal.Length	Sepal.Width	Petal.Length	Petal.Width
virginica	7.9	3.8	6.4	2.0
virginica	7.7	3.8	6.7	2.2
virginica	7.7	2.6	6.9	2.3
virginica	7.7	2.8	6.7	2.0
virginica	7.7	3.0	6.1	2.3
virginica	7.6	3.0	6.6	2.1
virginica	7.4	2.8	6.1	1.9
virginica	7.3	2.9	6.3	1.8
virginica	7.2	3.6	6.1	2.5
virginica	7.2	3.2	6.0	1.8

REconTools Function

REconTools Summarize over Tible

Use R4Econ's summary tool.

```r
df_summ_stats <- ff_summ_percentiles(tb_iris)
kable(t(df_summ_stats)) %>%
  kable_styling(bootstrap_options = c("striped", "hover", "responsive"))
```

stats	n	NAobs	ZEROobs	mean	sd	cv	min	p01	p05	p10	p25	p50
Petal.Length	150	0	0	3.758000	1.7652982	0.4697441	1.0	1.149	1.300	1.4	1.6	4.35
Petal.Width	150	0	0	1.199333	0.7622377	0.6355511	0.1	0.100	0.200	0.2	0.3	1.30
Sepal.Length	150	0	0	5.843333	0.8280661	0.1417113	4.3	4.400	4.600	4.8	5.1	5.80
Sepal.Width	150	0	0	3.057333	0.4358663	0.1425642	2.0	2.200	2.345	2.5	2.8	3.00