

# dplyr-and-tidyr-like functions written in base r

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## 1 Introduction

R-package m61r gathers functions similar to the ones present in dplyr and tidyr, but only written in base r, and without requiring any dependencies.  
All the functions only work with data.frames.

## 2 filter

```
> tmp <- filter_(CO2, ~Plant=="Qn1")
> head(tmp)

  Plant    Type Treatment conc uptake
1  Qn1 Quebec nonchilled   95   16.0
2  Qn1 Quebec nonchilled  175   30.4
3  Qn1 Quebec nonchilled  250   34.8
4  Qn1 Quebec nonchilled  350   37.2
5  Qn1 Quebec nonchilled  500   35.3
6  Qn1 Quebec nonchilled  675   39.2

>

> tmp <- filter_(CO2, ~Type=="Quebec")
> head(tmp)

  Plant    Type Treatment conc uptake
1  Qn1 Quebec nonchilled   95   16.0
2  Qn1 Quebec nonchilled  175   30.4
3  Qn1 Quebec nonchilled  250   34.8
4  Qn1 Quebec nonchilled  350   37.2
5  Qn1 Quebec nonchilled  500   35.3
6  Qn1 Quebec nonchilled  675   39.2

>
```

### 3 select

```
> tmp <- select_(CO2, ~Type)
> head(tmp)

      Type
1 Quebec
2 Quebec
3 Quebec
4 Quebec
5 Quebec
6 Quebec

>

> tmp <- select_(CO2, ~c(Plant,Type))
> head(tmp)

  Plant   Type
1 Qn1 Quebec
2 Qn1 Quebec
3 Qn1 Quebec
4 Qn1 Quebec
5 Qn1 Quebec
6 Qn1 Quebec

>

> tmp <- select_(CO2, ~-Type)
> head(tmp)

  Plant Treatment conc uptake
1 Qn1 nonchilled  95  16.0
2 Qn1 nonchilled 175  30.4
3 Qn1 nonchilled 250  34.8
4 Qn1 nonchilled 350  37.2
5 Qn1 nonchilled 500  35.3
6 Qn1 nonchilled 675  39.2

>

> tmp <- select_(CO2, variable=~-(Plant:Treatment))
> head(tmp)

  conc uptake
1    95    16.0
```

```

2 175 30.4
3 250 34.8
4 350 37.2
5 500 35.3
6 675 39.2

```

>

## 4 mutate/transmute

```

> tmp <- mutate_(CO2,z=^conc/uptake)
> head(tmp)

  Plant Type Treatment conc uptake      z
1  Qn1 Quebec nonchilled   95   16.0  5.937500
2  Qn1 Quebec nonchilled  175   30.4  5.756579
3  Qn1 Quebec nonchilled  250   34.8  7.183908
4  Qn1 Quebec nonchilled  350   37.2  9.408602
5  Qn1 Quebec nonchilled  500   35.3 14.164306
6  Qn1 Quebec nonchilled  675   39.2 17.219388

>

> tmp <- mutate_(CO2,mean=^mean(uptake))
> head(tmp)

  Plant Type Treatment conc uptake mean
1  Qn1 Quebec nonchilled   95   16.0    NA
2  Qn1 Quebec nonchilled  175   30.4    NA
3  Qn1 Quebec nonchilled  250   34.8    NA
4  Qn1 Quebec nonchilled  350   37.2    NA
5  Qn1 Quebec nonchilled  500   35.3    NA
6  Qn1 Quebec nonchilled  675   39.2    NA

>

> tmp <- mutate_(CO2,z1=^uptake/conc,y=^conc/100)
> head(tmp)

  Plant Type Treatment conc uptake      z1      y
1  Qn1 Quebec nonchilled   95   16.0  0.16842105  0.95
2  Qn1 Quebec nonchilled  175   30.4  0.17371429  1.75
3  Qn1 Quebec nonchilled  250   34.8  0.13920000  2.50
4  Qn1 Quebec nonchilled  350   37.2  0.10628571  3.50
5  Qn1 Quebec nonchilled  500   35.3  0.07060000  5.00
6  Qn1 Quebec nonchilled  675   39.2  0.05807407  6.75

```

```

>

> tmp <- transmute_(CO2,z2=~uptake/conc,y2=~conc/100)
> head(tmp)

      z2     y2
1 0.16842105 0.95
2 0.17371429 1.75
3 0.13920000 2.50
4 0.10628571 3.50
5 0.07060000 5.00
6 0.05807407 6.75

```

>

## 5 summarise

```

> tmp <- summarise_(CO2,mean=~mean(uptake),sd=~sd(uptake))
> tmp

      mean       sd
1 27.2131 10.81441

>

> tmp <- summarise_(CO2, group=~c(Type,Treatment),mean=~mean(uptake),sd=~sd(uptake))
> tmp

      Type Treatment     mean       sd
1    Quebec nonchilled 35.33333 9.596371
2    Quebec     chilled 25.95238 7.402136
3 Mississippi nonchilled 31.75238 9.644823
4 Mississippi     chilled 15.81429 4.058976

```

>

## 6 arrange/desange

```

> tmp <- arrange_(CO2,~c(conc))
> head(tmp)

      Plant   Type Treatment conc uptake
1    Qn1 Quebec nonchilled   95    16.0

```

```

2   Qn2 Quebec nonchilled  95   13.6
3   Qn3 Quebec nonchilled  95   16.2
4   Qc1 Quebec     chilled  95   14.2
5   Qc2 Quebec     chilled  95    9.3
6   Qc3 Quebec     chilled  95   15.1

>

> tmp <- arrange_(CO2, ~c(Treatment, conc, uptake))
> head(tmp)

  Plant      Type Treatment conc uptake
1 Mn1 Mississippi nonchilled  95   10.6
2 Mn3 Mississippi nonchilled  95   11.3
3 Mn2 Mississippi nonchilled  95   12.0
4 Qn2      Quebec nonchilled  95   13.6
5 Qn1      Quebec nonchilled  95   16.0
6 Qn3      Quebec nonchilled  95   16.2

>

> tmp <- desange_(CO2, ~c(Treatment, conc, uptake))
> head(tmp)

  Plant      Type Treatment conc uptake
1 Qc2      Quebec     chilled 1000   42.4
2 Qc3      Quebec     chilled 1000   41.4
3 Qc1      Quebec     chilled 1000   38.7
4 Mc1 Mississippi     chilled 1000   21.9
5 Mc3 Mississippi     chilled 1000   19.9
6 Mc2 Mississippi     chilled 1000   14.4

>

```

## 7 join

```

> authors <- data.frame(
+   surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),
+   nationality = c("US", "Australia", "US", "UK", "Australia"),
+   deceased = c("yes", rep("no", 4)))
> books <- data.frame(
+   name = I(c("Tukey", "Venables", "Tierney", "Ripley",
+             "Ripley", "McNeil", "R Core")),
+   title = c("Exploratory Data Analysis",
+            "Modern Applied Statistics ...",

```

```

+
+ "LISP-STAT",
+ "Spatial Statistics", "Stochastic Simulation",
+ "Interactive Data Analysis",
+ "An Introduction to R"),
+ other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))

```

## 7.1 inner join

```

> authors <- data.frame(
+   surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),
+   nationality = c("US", "Australia", "US", "UK", "Australia"),
+   deceased = c("yes", rep("no", 4)))
> books <- data.frame(
+   name = I(c("Tukey", "Venables", "Tierney", "Ripley",
+             "Ripley", "McNeil", "R Core")),
+   title = c("Exploratory Data Analysis",
+            "Modern Applied Statistics ...",
+            "LISP-STAT",
+            "Spatial Statistics", "Stochastic Simulation",
+            "Interactive Data Analysis",
+            "An Introduction to R"),
+   other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))
> tmp <- inner_join_(authors, books, by.x = "surname", by.y = "name")
> tmp

```

	surname	nationality	deceased	title	other.author
1	McNeil	Australia	no	Interactive Data Analysis	<NA>
2	Ripley	UK	no	Spatial Statistics	<NA>
3	Ripley	UK	no	Stochastic Simulation	<NA>
4	Tierney	US	no	LISP-STAT	<NA>
5	Tukey	US	yes	Exploratory Data Analysis	<NA>
6	Venables	Australia	no	Modern Applied Statistics ...	Ripley

>

## 7.2 left join

```

> authors <- data.frame(
+   surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),
+   nationality = c("US", "Australia", "US", "UK", "Australia"),
+   deceased = c("yes", rep("no", 4)))
> books <- data.frame(
+   name = I(c("Tukey", "Venables", "Tierney", "Ripley",
+             "Ripley", "McNeil", "R Core")),
+   title = c("Exploratory Data Analysis",
+            "Modern Applied Statistics ...",
+            "LISP-STAT",
+            "Spatial Statistics", "Stochastic Simulation",
+            "Interactive Data Analysis",
+            "An Introduction to R"),
+   other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))

```

```

+
+           "Spatial Statistics", "Stochastic Simulation",
+           "Interactive Data Analysis",
+           "An Introduction to R"),
+           other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))
> tmp <- left_join_(authors, books, by.x = "surname", by.y = "name")
> tmp

  surname nationality deceased          title other.author
1   McNeil     Australia      no  Interactive Data Analysis      <NA>
2   Ripley        UK      no       Spatial Statistics      <NA>
3   Ripley        UK      no  Stochastic Simulation      <NA>
4  Tierney        US      no        LISP-STAT      <NA>
5   Tukey        US     yes Exploratory Data Analysis      <NA>
6 Venables     Australia     no Modern Applied Statistics ...        Ripley

>

```

### 7.3 right join

```

> authors <- data.frame(
+   surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),
+   nationality = c("US", "Australia", "US", "UK", "Australia"),
+   deceased = c("yes", rep("no", 4)))
> books <- data.frame(
+   name = I(c("Tukey", "Venables", "Tierney", "Ripley",
+             "R Core", "McNeil", "R Core")),
+   title = c("Exploratory Data Analysis",
+             "Modern Applied Statistics ...",
+             "LISP-STAT",
+             "Spatial Statistics", "Stochastic Simulation",
+             "Interactive Data Analysis",
+             "An Introduction to R"),
+   other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))
> tmp <- right_join_(authors, books, by.x = "surname", by.y = "name")
> tmp

  surname nationality deceased          title other.author
1   McNeil     Australia      no  Interactive Data Analysis      <NA>
2   R Core        <NA>    <NA> An Introduction to R Venables & Smith
3   Ripley        UK      no       Spatial Statistics      <NA>
4   Ripley        UK      no  Stochastic Simulation      <NA>
5  Tierney        US      no        LISP-STAT      <NA>
6   Tukey        US     yes Exploratory Data Analysis      <NA>
7 Venables     Australia     no Modern Applied Statistics ...        Ripley

>

```

## 7.4 full join

```
> authors <- data.frame(  
+     surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),  
+     nationality = c("US", "Australia", "US", "UK", "Australia"),  
+     deceased = c("yes", rep("no", 4)))  
> books <- data.frame(  
+     name = I(c("Tukey", "Venables", "Tierney", "Ripley",  
+                 "Ripley", "McNeil", "R Core")),  
+     title = c("Exploratory Data Analysis",  
+               "Modern Applied Statistics ...",  
+               "LISP-STAT",  
+               "Spatial Statistics", "Stochastic Simulation",  
+               "Interactive Data Analysis",  
+               "An Introduction to R"),  
+     other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))  
> tmp <- full_join_(authors, books, by.x = "surname", by.y = "name")  
> tmp  
  
surname nationality deceased          title      other.author  
1   McNeil    Australia      no  Interactive Data Analysis             <NA>  
2   R Core        <NA>      <NA>  An Introduction to R Venables & Smith  
3   Ripley       UK        no   Spatial Statistics             <NA>  
4   Ripley       UK        no   Stochastic Simulation             <NA>  
5  Tierney       US        no      LISP-STAT             <NA>  
6   Tukey        US       yes Exploratory Data Analysis             <NA>  
7  Venables    Australia      no Modern Applied Statistics ...        Ripley  
  
>
```

## 7.5 semi join

```
> authors <- data.frame(  
+     surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),  
+     nationality = c("US", "Australia", "US", "UK", "Australia"),  
+     deceased = c("yes", rep("no", 4)))  
> books <- data.frame(  
+     name = I(c("Tukey", "Venables", "Tierney", "Ripley",  
+                 "Ripley", "McNeil", "R Core")),  
+     title = c("Exploratory Data Analysis",  
+               "Modern Applied Statistics ...",  
+               "LISP-STAT",  
+               "Spatial Statistics", "Stochastic Simulation",  
+               "Interactive Data Analysis",  
+               "An Introduction to R"),  
+     other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))  
> tmp <- semi_join_(authors, books, by.x = "surname", by.y = "name")  
> tmp
```

```

surname nationality deceased
1 Tukey US yes
2 Venables Australia no
3 Tierney US no
4 Ripley UK no
5 McNeil Australia no

```

>

## 7.6 anti join

```

> authors <- data.frame(
+   surname = I(c("Tukey", "Venables", "Tierney", "Ripley", "McNeil")),
+   nationality = c("US", "Australia", "US", "UK", "Australia"),
+   deceased = c("yes", rep("no", 4)))
> books <- data.frame(
+   name = I(c("Tukey", "Venables", "Tierney", "Ripley",
+             "Ripley", "McNeil", "R Core")),
+   title = c("Exploratory Data Analysis",
+             "Modern Applied Statistics ...",
+             "LISP-STAT",
+             "Spatial Statistics", "Stochastic Simulation",
+             "Interactive Data Analysis",
+             "An Introduction to R"),
+   other.author = c(NA, "Ripley", NA, NA, NA, NA, "Venables & Smith"))
> tmp <- anti_join_(authors, books, by.x = "surname", by.y = "name")
> tmp

```

```
[1] surname      nationality deceased
<0 rows> (or 0-length row.names)
```

```

> tmp <- anti_join_(books, authors, by.x = "name", by.y = "surname")
> tmp

```

```

name          title      other.author
7 R Core An Introduction to R Venables & Smith

```

>

## 8 reshape: merge/spread

### 8.1 merge

```

> df3 <- data.frame(id = 1:4,
+                     age = c(40,50,60,50),

```

```

+
+           dose.a1 = c(1,2,1,2),
+           dose.a2 = c(2,1,2,1),
+           dose.a14 = c(3,3,3,3))
> df3

   id age dose.a1 dose.a2 dose.a14
1  1  40      1      2      3
2  2  50      2      1      3
3  3  60      1      2      3
4  4  50      2      1      3

> gather_(df3,pivot = c("id","age"))

  id age parameters values
1  1  40      dose.a1     1
2  2  50      dose.a1     2
3  3  60      dose.a1     1
4  4  50      dose.a1     2
5  1  40      dose.a2     2
6  2  50      dose.a2     1
7  3  60      dose.a2     2
8  4  50      dose.a2     1
9  1  40      dose.a14    3
10 2  50      dose.a14    3
11 3  60      dose.a14    3
12 4  50      dose.a14    3

>

```

## 8.2 spread

```

> df3 <- data.frame(id = 1:4,
+                      age = c(40,50,60,50),
+                      dose.a1 = c(1,2,1,2),
+                      dose.a2 = c(2,1,2,1),
+                      dose.a14 = c(3,3,3,3))
> df3

   id age dose.a1 dose.a2 dose.a14
1  1  40      1      2      3
2  2  50      2      1      3
3  3  60      1      2      3
4  4  50      2      1      3

> gather_(df3,pivot = c("id","age"))

```

```

      id age parameters values
1   1  40    dose.a1     1
2   2  50    dose.a1     2
3   3  60    dose.a1     1
4   4  50    dose.a1     2
5   1  40    dose.a2     2
6   2  50    dose.a2     1
7   3  60    dose.a2     2
8   4  50    dose.a2     1
9   1  40    dose.a14    3
10  2  50    dose.a14    3
11  3  60    dose.a14    3
12  4  50    dose.a14    3

> df4 <- gather_(df3,pivot = c("id","age"))
> df5 <- rbind(df4,
+   data.frame(id=5, age=20,parameters="dose.a14",values=8),
+   data.frame(id=6, age=10,parameters="dose.a1",values=5))
> df5

      id age parameters values
1   1  40    dose.a1     1
2   2  50    dose.a1     2
3   3  60    dose.a1     1
4   4  50    dose.a1     2
5   1  40    dose.a2     2
6   2  50    dose.a2     1
7   3  60    dose.a2     2
8   4  50    dose.a2     1
9   1  40    dose.a14    3
10  2  50    dose.a14    3
11  3  60    dose.a14    3
12  4  50    dose.a14    3
13  5  20    dose.a14    8
14  6  10    dose.a1     5

> spread_(df5,col_name="parameters",col_values="values",pivot=c("id","age"))

      id age dose.a1 dose.a14 dose.a2
1   1  40      1      3      2
2   2  50      2      3      1
3   3  60      1      3      2
4   4  50      2      3      1
5   5  20     NA      8     NA
6   6  10      5     NA     NA

```