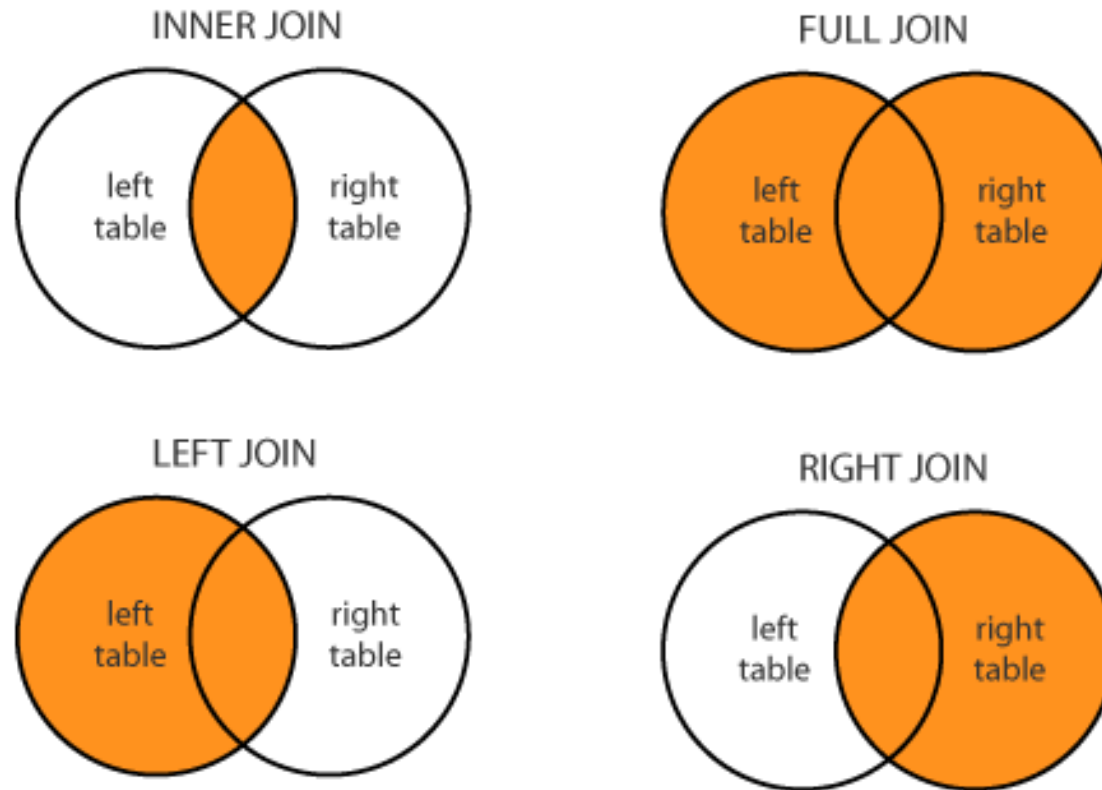




# DATAFRAMES PART 2

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# JOINING DATSETS



<https://i.stack.imgur.com/iJUMl.png>

```
# Inner join  
merge(set1, set2)
```

```
# Outer join  
  
merge(x = set1, y = set2,  
      by = "IdClient", all =  
      TRUE)
```

```
# Left outer join  
  
merge(x = set1, y = set2,  
      by = "IdClient", all.x =  
      TRUE)
```

```
# Right outer join  
  
merge(x = set1, y = set2,  
      by = "IdKlienta", all.y =  
      TRUE)
```

# TEXT DATA — USEFUL FUNCTIONS

„Cutting-off” signs at given position:

- `substr()`

Merging elements into one text sequence:

- `paste()` `paste0()`

Returning the length of elements in subsequent positions of text vector:

- `nchar()`

Changing letter capitalization:

- `toupper()` `tolower()`

# ORDERING DATA

Transpose data/matrix:

- `t()`

Sorting a variable by its values:

- `sort()`

Ordering dataset by values of given variable:

- `order()`

# FILTERING DATA

Choosing rows by the condition on column values:

- `set12[set1$Product == "Bike", ]`

Creating a subset using certain feature:

- `subset(set12, Product == "Bike")`

# CHECKING FOR MISSING DATA

Checking which values are missing:

- `is.na()`

Checking which rows do not include missing values:

- `complete.cases()`

# RANDOM NUMBERS

Random number generator:

- `runif()`

Specifying the starting point  
for random number generating:

- `set.seed()`

# USING PACKAGES

Using a package in R:

1. install the package when using it for the first time in the given R version on a given computer

```
install.packages("packageName")
```

2. load the package every time you'd like to use it - once per working R session

```
library("packageName")
```

```
library(packageName) #both notations are ok
```



# Package documentation website

cluster: "Finding Groups in Data": Cluster Analysis Extended Rousseeuw et al.

Methods for Cluster analysis. Much extended the original from Peter Rousseeuw, Anja Struyf and Mia Hubert, bas

Version: 2.1.4 **Pacversion ersion**

Priority: recommended



Depends: R ( $\geq 3.5.0$ )

Imports: graphics, grDevices, stats, utils

Suggests: [MASS](#), [Matrix](#)

**Dependencies**

Published: 2022-08-22

Author: Martin Maechler  [aut, cre], Peter Rousseeuw  [aut] (Fortran original), Anja Struyf [au  
(volume.ellipsoid({d >= 3}))]

**Authors and maintainers**

Maintainer: Martin Maechler <maechler at stat.math.ethz.ch>

License: [GPL-2](#) | [GPL-3](#) [expanded from: GPL ( $\geq 2$ )]

URL: <https://svn.r-project.org/R-packages/trunk/cluster/>

**Webpage of the package – usually  
with tutorials and further descriptions**

NeedsCompilation: yes

Citation: [cluster citation info](#)

**Citation**

Materials: [README](#) [NEWS](#) [ChangeLog](#)

In views: [Cluster](#), [Environmetrics](#), [Robust](#)

CRAN checks: [cluster results](#)

Documentation:

Reference manual: [cluster.pdf](#) **Documentation!!!**

Downloads:

Package source: [cluster\\_2.1.4.tar.gz](#) **Source code**

Windows binaries: r-devel: [cluster\\_2.1.4.zip](#), r-release: [cluster\\_2.1.4.zip](#), r-oldrel: [cluster\\_2.1.4.zip](#)

macOS binaries: r-release (arm64): [cluster\\_2.1.4.tgz](#), r-oldrel (arm64): [cluster\\_2.1.4.tgz](#), r-release (x86\_64): [clu](#)

Old sources: [cluster archive](#)

**Older versions**

Reverse dependencies:

Reverse depends: [abodOutlier](#), [aCGH](#), [AssocTests](#), [bios2mds](#), [BoutrosLab](#), [plotting.general](#), [briKmeans](#), [ClassDis](#),  
[NMF](#), [nomclust](#), [optpart](#), [Oscope](#), [PAMhm](#), [pamr](#), [RnBeads](#), [RPM](#), [saccadr](#), [STROMA4](#), [Stru](#)

Reverse imports: [ADaCGH2](#), [adiv](#), [ADPclust](#), [adSplit](#), [agricolae](#), [anocva](#), [Anthropometry](#), [aPCoA](#), [aqp](#), [artMS](#), [A](#)  
[clusterExperiment](#), [clusterHD](#), [Clustering](#), [clustrd](#), [CLUSTShiny](#), [cobiclust](#), [coca](#), [CoGAPS](#), [co](#)  
[EvaluateCore](#), [EvoPhylo](#), [factoextra](#), [FactoMineR](#), [FADPclust](#), [FairMclus](#), [fdm2id](#), [flowStats](#), [f](#)

## R Updating Loaded Packages



One or more of the packages to be updated are currently loaded. Restarting R prior to install is highly recommended.

RStudio can restart R before installing the requested packages. All work and data will be preserved during restart.

Do you want to restart R prior to install?

Yes

No

Cancel



If you try to install a package that is already loaded to the R session – and it's outdated compared to the newest version available on CRAN:

CANCEL → You will just close this pop-up and stay with the currently loaded package version.

NO → You will reinstall the package to the newest version, then you need to reload it again. R session won't be restarted – there may be some issues with the package running.

YES → You will reinstall the package to the newest version, then you need to reload it again. R session WILL BE RESTARTED. R Studio tries to preserve your work and keep it in the memory but you should be prepared to re-run all your code. Stable solution – updated package will be working as it should.