

# Learning to Reorder Factor Levels in R: A Comprehensive Guide with Examples

Authored by  
**Mohammed loot**

November 5, 2025

## RECOMMENDED CITATION

Mohammed loot (2025). *Learning to Reorder Factor Levels in R: A Comprehensive Guide with Examples*. PSYCHOLOGICAL STATISTICS. Retrieved from <https://statistics.arabpsychology.com/?p=10974>

## Introduction to Factors and Ordering in R

When conducting statistical analysis and data manipulation within the [R programming language](#), handling [categorical data](#) is a frequent and crucial task. R utilizes a specialized data structure known as the [factor](#) to efficiently store and manage these variables. Factors are essential for almost all modeling and visualization operations in R because they treat character strings not merely as text, but as predefined categories with specific, discrete values.

A factor is fundamentally defined by its underlying values and its set of unique identifiers, referred to as **levels**. By default, R automatically assigns these levels either based on the order of their appearance in the dataset or, more commonly, in standard alphabetical (lexicographical) order. While this automatic assignment is convenient, it rarely aligns with the specific analytical requirements of a project. Consider, for example, survey responses where the logical progression is "Poor," "Fair," "Good," and "Excellent." Alphabetical sorting would incorrectly sequence these as "Excellent," "Fair," "Good," "Poor."

The need for a custom arrangement--an order that reflects logical importance, temporal sequence, or specific reporting standards--makes manual reordering of **factor levels** indispensable. Without this control, visualizations and statistical outputs may be misleading or fail to conform to expected standards. This tutorial will provide a robust, reliable method using base R functions to take complete control over how your categorical variables are sequenced and interpreted, ensuring precision in your data presentation.

### The Core Method: Utilizing the `factor()` Function

The fundamental approach to redefining the sequence of levels within an [R factor](#) involves reapplying the built-in `factor()` function to the variable itself. This method is considered the standard base R practice because it is explicit, reliable, and does not require external packages. The key to successful reordering lies in utilizing the `levels` argument, where you provide a vector specifying the complete, newly desired arrangement.

When you use the `factor()` function in this manner, R essentially rebuilds the factor structure. It maps the existing data values to the new internal level definitions, effectively overwriting the previous order while maintaining the integrity of the underlying data points. This process ensures that subsequent operations, such as filtering, grouping, or plotting, respect the new, customized sequence.

It is absolutely **crucial** to understand the dependency of this method: the vector supplied to the `levels` argument must contain every unique level currently present in the factor. If any existing level is omitted from the new vector, R will convert all observations corresponding to that missing level into [NA](#) (Not Available). This silent conversion can lead to data loss or incorrect analysis,

making careful verification of the level list a mandatory prerequisite for reordering.

The general structure for this core operation, where the complete sequence is explicitly defined, is presented below, illustrating how the new level arrangement is passed to the function:

```
factor_variable <- factor(factor_variable, levels=c('this', 'that', 'those', ...))
```

Mastering this precise syntax provides the foundation for effective management of categorical variables in R. We now transition to a hands-on example to demonstrate how this technique is applied within the common context of an R [data frame](#).

## Practical Demonstration: Preparing the Data Frame

To clearly illustrate the process of custom level reordering, we will begin by constructing a simple, representative R object: a [data frame](#). This structure, which we will name `df`, is the most common container for analytical data in R and provides a realistic environment for factor manipulation. Our data frame will include two variables: `region`, which will serve as our target categorical variable, and `sales`, a corresponding quantitative measure.

In this example, the `region` variable is intentionally initialized as a **factor** immediately upon creation. This reflects a typical scenario often encountered after importing data, where character vectors representing categories are automatically coerced into factors by R's read functions. Currently, the levels will be ordered alphabetically based on the input sequence (A, B, C, D, E).

The following code block details the creation of this sample data structure in R, followed by the output displaying the data frame contents:

```
#create data frame  
df <- data.frame(region=factor(c('A', 'B', 'C', 'D', 'E')),  
sales=c(12, 18, 21, 14, 34))
```

```
#view data frame
```

```
df
```

```
region sales
```

```
1 A 12
```

```
2 B 18
```

```
3 C 21
```

```
4 D 14
```

```
5 E 34
```

Prior to applying any changes, the best practice is to always inspect the current, default sequence of the factor levels. We can quickly confirm the initial ordering by applying the standard `levels()` function specifically to the `region` column within our data frame. This reveals R's current internal definition of the categories:

```
#display factor levels for region  
levels(df$region)
```

```
"A" "B" "C" "D" "E"
```

As confirmed by the output, the initial level order is the basic alphabetical sequence. Our objective in the subsequent step is to deliberately disrupt this default order and impose a completely customized arrangement that better suits hypothetical reporting needs.

## Implementing Custom Level Reordering

Imagine that for a specific monthly business report, the regions must be presented in a non-alphabetical sequence reflecting analytical priority, such as sales performance or geographical importance. Let us define the required custom order as: A, E, D, C, B. This sequence deviates entirely from the standard lexicographical sort, necessitating a manual factor definition update.

The procedure involves reapplying the `factor()` function to the `df$region` column, passing the explicitly defined vector `c('A', 'E', 'D', 'C', 'B')` to the `levels` argument. This critical assignment step ensures that the old factor definition is replaced by the new one, immediately governing how R treats this column in all subsequent operations.

The following implementation code executes the reordering and verifies the resulting structure change by re-checking the levels:

```
#re-order factor levels for region  
df$region <- factor(df$region, levels=c('A', 'E', 'D', 'C', 'B'))  
  
#display factor levels for region  
levels(df$region)
```

```
"A" "E" "D" "C" "B"
```

The output confirms the successful manipulation. The factor levels for `region` are now precisely arranged in the sequence "A", "E", "D", "C", "B". It is essential to remember that while the raw sales data remains paired correctly with its region label, the categorical structure now dictates this custom sequence for sorting and aggregation. This successful redefinition is the prerequisite for

generating accurate, custom-ordered visualizations.

## Application: Controlling Data Visualization Output

The primary motivation for manually reordering factor levels is to gain absolute control over the presentation of categorical data in plots. When R generates visualizations involving categories--such as [barplots](#), box plots, or grouped scatterplots--the sequence in which the elements appear along the axis is directly determined by the internal ordering of the factor levels. If the levels are not correctly ordered, the visual narrative may be fundamentally flawed.

To ensure that a visualization, such as a barplot comparing sales across regions, strictly adheres to our newly defined sequence (A, E, D, C, B), we must often take an extra step: we must sort the rows of the data frame itself based on the factor level order. Although modern plotting libraries often respect factor order automatically, explicitly sorting the data frame rows guarantees the desired arrangement, especially when utilizing base R plotting functions like `barplot()`.

We achieve this sorting by combining the `order()` function with the current factor levels of the `region` variable. Once the data frame is properly sequenced, we proceed to generate the barplot, ensuring the bars are placed in our custom order:

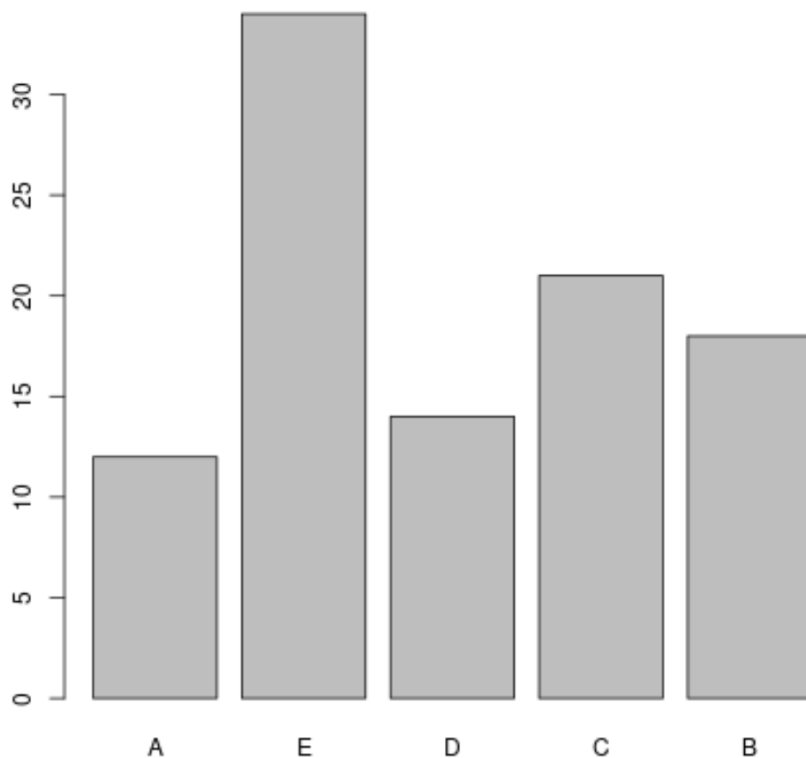
```
#re-order data frame based on factor levels for region
```

```
df <- df
```

```
#create barplot and place bars in order based on factor levels for region
```

```
barplot(df$sales, names=df$region)
```

The resulting visual output now perfectly reflects the analytical requirement. The bars representing the sales figures are presented in the sequence A, E, D, C, B, rather than the default alphabetical order. This ability to link the data's internal categorical definition directly to its visual representation is a cornerstone of effective and professional data reporting.



A quick visual inspection confirms that the bars align with the customized order of the **factor levels**. This explicit control over sequence is necessary when communicating complex data relationships clearly and logically.

## Streamlining Factor Manipulation with the Tidyverse

While the base R method using `factor(..., levels=...)` is foundational and robust across all R environments, modern data science workflows frequently benefit from specialized packages that simplify factor management. The **tidyverse** collection, specifically the dedicated [forcats](#) package, provides a suite of highly intuitive functions designed to overcome the common frustrations associated with traditional factor handling.

For tasks involving manual, explicit reordering, `forcats` offers the streamlined function `fct_relevel()`. This function improves readability and reduces potential errors because it allows the user to specify only the levels they wish to move to the front or back of the sequence, rather than requiring a complete list of all existing levels. For instance, if you only wanted to move regions "C" and "B" to the beginning, the syntax is significantly cleaner: `df$region <- fct_relevel(df$region, "C", "B")`. All other levels automatically retain their relative order.

Furthermore, `forcats` excels in scenarios requiring statistical reordering--where the level order is determined dynamically by another variable in the dataset. If the requirement was to order the

regions based on their actual corresponding sales figures (the values in the `sales` column), the function `fct_reorder()` would be the ideal choice. This powerful tool dynamically arranges the factor levels based on an aggregation or summary statistic, making it indispensable for creating bar charts where the bars must be sequenced by magnitude (e.g., from highest to lowest sales volume).

Choosing between the base R approach and the Tidyverse tools depends heavily on the complexity and volume of the factor manipulation required. For simple, one-off reordering, the base R method is perfectly adequate. However, for advanced, statistically driven ordering, or for integrating factor management into efficient data pipelines using the pipe operator, `forcats` provides vastly superior efficiency, safety, and code clarity.

## Conclusion and Summary of Best Practices

The ability to accurately and manually reorder **factor levels** is an essential skill for any R user involved in data preparation, statistical modeling, or creating high-quality data visualizations. While R's default alphabetical ordering serves as a starting point, custom orders are frequently mandated by business logic, statistical requirements, or the need to present a clear analytical narrative.

The primary, robust technique for achieving this customization involves redefining the factor variable using the base R `factor()` function, paired with the explicit definition of the level sequence via the `levels` argument. This ensures that the data structure is updated without compromising the integrity of the underlying observations.

To ensure successful factor management in all future projects, keep these key best practices in mind:

**Completeness is Critical:** Always confirm that the vector provided to the `levels` argument includes every unique category present in the factor. Failure to do so will result in existing data points being silently converted to missing values ([NA](#)).

**Visualization Control:** Understand that reordering factor levels is the direct, explicit mechanism for dictating the sequence of elements in [barplots](#) and other plots that rely on categorical axes.

**Efficiency through Specialization:** For statistical reordering (e.g., sorting by mean value) or complex manipulation in a pipeline, leverage specialized tools such as the `forcats` package for enhanced readability and fewer manual errors.

By mastering factor level reordering, you guarantee that your categorical data is not only analyzed correctly but also presented in a manner that maximizes clarity and aligns perfectly with the intended interpretation.

Explore more **R tutorials** to enhance your skills in analyzing and manipulating data structures efficiently.