

# C

## Compendium

### Elementary

#### Commands

<code>ls()</code> or <code>objects()</code>	List objects in workspace
<code>rm(object)</code>	Delete object
<code>search()</code>	Search path

#### Variable names

Combinations of letters, digits and period. Must not start with a digit.  
Avoid starting with period.

#### Assignments

<code>&lt;-</code>	Assign value to variable
<code>-&gt;</code>	Assignment “to the right”
<code>&lt;&lt;-</code>	Global assignment (in functions)

## Operators

### Arithmetic

<code>+</code>	Addition
<code>-</code>	Subtraction, sign
<code>*</code>	Multiplication
<code>/</code>	Division
<code>^</code>	Raise to power
<code>% / %</code>	Integer division
<code>%%</code>	Remainder from integer division

### Logical and relational

<code>==</code>	Equal to
<code>!=</code>	Not equal to
<code>&lt;</code>	Less than
<code>&gt;</code>	Greater than
<code>&lt;=</code>	Less than or equal to
<code>&gt;=</code>	Greater than or equal to
<code>is.na(x)</code>	Missing?
<code>&amp;</code>	Logical AND
<code> </code>	Logical OR
<code>!</code>	Logical NOT

`&` and `|` are *elementwise*. See “Programming” (p. 258) for `&&` and `||`.

## Vectors and data types

### Generating

<code>numeric(25)</code>	25 zeros
<code>character(25)</code>	25 × "
<code>logical(25)</code>	25 × FALSE
<code>seq(-4, 4, 0.1)</code>	Sequence: -4.0 -3.9 3.8 ... 3.9 4.0
<code>1:10</code>	Same as <code>seq(1, 10, 1)</code>
<code>c(5, 7, 9, 13, 1:5)</code>	Concatenation: 5 7 9 13 1 2 3 4 5
<code>rep(1, 10)</code>	1 1 1 1 1 1 1 1 1 1
<code>g1(3, 2, 12)</code>	Factor with 3 levels, repeat each level in blocks of 2, up to length 12 (i.e., 1 1 2 2 3 3 1 1 2 2 3 3)

### Coercion

<code>as.numeric(x)</code>	Convert to numeric
<code>as.character(x)</code>	Convert to text string
<code>as.logical(x)</code>	Convert to logical
<code>factor(x)</code>	Create factor from vector x

Re. factors, see also "Tabulation, grouping, recoding" (p. 253).

## Data frames

<code>data.frame(height = c(165, 185), weight = c(90, 65))</code>	Data frame with two named vectors
<code>data.frame(height, weight)</code>	Collect vectors into data frame
<code>dfr\$var</code>	Select vector var in data frame dfr
<code>attach(dfr)</code>	Put data frame in search path
<code>detach()</code>	— and remove it from path

Attached data frames always come *after* `.GlobalEnv` in the search path.

Attached data frames are copies; subsequent changes to `dfr` have no effect.

## Numerical functions

### Mathematical

<code>log(x)</code>	Logarithm of $x$ , natural (base- $e$ ) logarithm
<code>log10(x)</code>	Base-10 logarithm
<code>exp(x)</code>	Exponential function $e^x$
<code>sin(x)</code>	Sine
<code>cos(x)</code>	Cosine
<code>tan(x)</code>	Tangent
<code>asin(x)</code>	Arcsin (inverse sine)
<code>acos(x)</code>	
<code>atan(x)</code>	
<code>min(x)</code>	Smallest value in vector
<code>min(x1, x2, ...)</code>	minimum over several vectors (one number)
<code>max(x)</code>	Largest value in vector
<code>range(x)</code>	Like $c(\min(x), \max(x))$
<code>pmin(x1, x2, ...)</code>	Parallel (elementwise) minimum over multiple equally long vectors
<code>pmax(x1, x2, ...)</code>	Parallel maximum
<code>length(x)</code>	Number of elements in vector
<code>sum(complete.cases(x))</code>	Number of non-missing elements in vector

### Statistical

<code>mean(x)</code>	Average
<code>sd(x)</code>	Standard deviation
<code>var(x)</code>	Variance
<code>median(x)</code>	Median
<code>quantile(x, p)</code>	Quantiles
<code>cor(x, y)</code>	Correlation

## Indexing/selection

<code>x[1]</code>	First element
<code>x[1:5]</code>	Subvector containing first five elements
<code>x[c(2,3,5,7,11)]</code>	Element nos. 2, 3, 5, 7, and 11
<code>x[y&lt;=30]</code>	Selection by logical expression
<code>x[sex=="male"]</code>	Selection by factor variable
<code>i &lt;- c(2,3,5,7,11); x[i]</code>	Selection by numerical variable
<code>l &lt;- (y&lt;=30); x[l]</code>	Selection by logical variable

## Matrices and data frames

<code>m[4, ]</code>	Fourth row
<code>m[, 3]</code>	Third column
<code>dfr[dfr\$var&lt;=30, ]</code>	Partial data frame
<code>subset(dfr, var&lt;=30)</code>	Same, often simpler

## Input of data

<code>data(name)</code>	Built-in data set
<code>read.table("filename")</code>	Read from external file

## Common arguments to `read.table`

<code>header=TRUE</code>	First line has variable names
<code>sep = " , "</code>	Data are separated by commas
<code>dec = " , "</code>	Decimal point is comma
<code>na.strings=". "</code>	Missing value is dot

## Variants of `read.table`

<code>read.csv("filename")</code>	Comma separated
<code>read.delim("filename")</code>	Tab delimited
<code>read.csv2("filename")</code>	Semicolon separated, comma decimal point
<code>read.delim2("filename")</code>	Tab delimited, comma decimal point

These all set `header=TRUE`

## Missing values

### Functions

<code>is.na(x)</code>	Logical vector. TRUE where x has NA
<code>complete.cases(x1, x2, ...)</code>	Neither missing in x1, nor x2, nor...

### Arguments to other functions

<code>na.rm=</code>	In statistical functions. Remove missing if TRUE, return NA if FALSE.
<code>na.last=</code>	In sort TRUE, FALSE and NA means, respectively, "last", "first", and "throw away".
<code>na.action=</code>	In lm, etc., values na.fail, na.omit, na.exclude. Also in options ("na.action").
<code>na.print=</code>	In summary and print.default: How to represent NA in output.
<code>na.strings=</code>	In read.table(): Code(s) for NA in input.

## Tabulation, grouping, recoding

<code>table(f1, ...)</code>	(Cross)tabulation
<code>tapply(x, f, mean)</code>	Table of means
<code>factor(x)</code>	Convert vector to factor
<code>cut(x, breaks)</code>	Groups from cutpoints for continuous variable

### Arguments to `factor`

<code>levels</code>	Values of $x$ to code. Use if some values are not present in data, or if the order would be wrong.
<code>labels</code>	Values associated with factor levels.
<code>exclude</code>	Values to exclude. Default NA. Set to NULL to have missing values included as a level.

### Arguments to `cut`

<code>breaks</code>	Cutpoints. Note values of $x$ outside <code>breaks</code> gives NA. Can also be a single number, the number of cutpoints (not very useful).
<code>labels</code>	Names for groups. Default is $(0, 30]$ , etc.
<code>right</code>	Right endpoint included? (FALSE: left)

### Recoding factors

<code>levels(f) &lt;- names</code>	New level names
<code>factor(newcodes[f])</code>	Combining levels: <code>newcodes</code> , e.g., <code>c(1, 1, 1, 2, 3)</code> to amalgamate the first 3 of 5 groups.

## Statistical distributions

### Normal distribution

<code>dnorm(x)</code>	Density
<code>pnorm(x)</code>	Cumulative distribution function, $P(X \leq x)$
<code>qnorm(p)</code>	$p$ -quantile, $x : P(X \leq x) = p$
<code>rnorm(n)</code>	n (pseudo-)random normally distributed numbers

### Distributions

<code>pnorm(x, mean, sd)</code>	Normal
<code>plnorm(x, mean, sd)</code>	Lognormal
<code>pt(x, df)</code>	Student's $t$
<code>pf(x, n1, n2)</code>	$F$ distribution
<code>pchisq(x, df)</code>	$\chi^2$
<code>pbinom(x, n, p)</code>	Binomial
<code>ppois(x, lambda)</code>	Poisson
<code>punif(x, min, max)</code>	Uniform
<code>pexp(x, rate)</code>	Exponential
<code>pgamma(x, shape, scale)</code>	Gamma
<code>pbeta(x, a, b)</code>	Beta

Same convention (d-q-r) for density, quantiles, and random numbers as for normal distribution.

## Statistical standard methods

### Continuous response

<code>t.test</code>	One- and two-sample <i>t</i> test
<code>pairwise.t.test</code>	Pairwise comparisons
<code>cor.test</code>	Correlation
<code>var.test</code>	Comparison of two variances ( <i>F</i> test)
<code>lm(y ~ x)</code>	Regression analysis
<code>lm(y ~ f)</code>	One-way analysis of variance
<code>lm(y ~ f1 + f2)</code>	Two-way analysis of variance
<code>lm(y ~ f + x)</code>	Analysis of covariance
<code>lm(y ~ x1 + x2 + x3)</code>	Multiple regression analysis
<code>bartlett.test</code>	Bartlett's test ( <i>k</i> variances)
<b>Nonparametric:</b>	
<code>wilcox.test</code>	One- and two-sample Wilcoxon test
<code>kruskal.test</code>	Kruskal–Wallis test
<code>friedman.test</code>	Friedman's two-way analysis of variance
<b>cor.test variants:</b>	
<code>method = "kendall"</code>	Kendall's $\tau$
<code>method = "spearman"</code>	Spearman's $\rho$

### Discrete response

<code>binom.test</code>	Binomial test (incl. sign test)
<code>prop.test</code>	Comparison of proportions
<code>prop.trend.test</code>	Test for trend in relative proportions
<code>fisher.test</code>	Exact test in small tables
<code>chisq.test</code>	$\chi^2$ test
<code>glm(y ~ x1+x2+x3, binomial)</code>	Logistic regression

## Models

### Model formulas

$\sim$	Described by
$+$	Additive effects
$:$	Interaction
$*$	Main effects + interaction ( $a * b = a + b + a:b$ )
$-1$	Remove intercept

Classifications are represented by descriptive variable being a *factor*.

### Linear and generalized linear models

<code>lm.out &lt;- lm(y ~ x)</code>	Fit model and save result
<code>summary(lm.out)</code>	Coefficients, etc.
<code>anova(lm.out)</code>	Analysis of variance table
<code>fitted(lm.out)</code>	Fitted values
<code>resid(lm.out)</code>	Residuals
<code>predict(lm.out, newdata)</code>	Predictions for new data frame
<code>glm(y ~ x, binomial)</code>	Logistic regression

### Diagnostics

<code>rstudent(lm.out)</code>	Studentized residuals
<code>dfbetas(lm.out)</code>	Change in $\beta$ if obs. removed
<code>dffits(lm.out)</code>	Change in fit if obs. removed

### Survival analysis

<code>S &lt;- Surv(time, ev)</code>	Create survival object
<code>survfit(S)</code>	Kaplan-Meier estimate
<code>plot(survfit(S))</code>	Survival curve
<code>survdiff(S ~ g)</code>	(Log-rank) test for equal survival curves
<code>coxph(S ~ x1 + x2)</code>	Cox's proportional hazards model

## Graphics

### Standard plots

<code>plot()</code>	Scatterplot (and more)
<code>hist()</code>	Histogram
<code>boxplot()</code>	Box-and-whiskers plot
<code>stripplot()</code>	Stripplot
<code>barplot()</code>	Bar diagram
<code>dotplot()</code>	Dot diagram
<code>piechart()</code>	Cakes...
<code>interaction.plot()</code>	Interaction plot

### Plotting elements

<code>lines()</code>	Lines
<code>abline()</code>	Line given by intercept and slope (and more)
<code>points()</code>	Points
<code>segments()</code>	Line segments
<code>arrows()</code>	Arrows (NB: angle=90 for error bars)
<code>axis()</code>	Axis
<code>box()</code>	Frame around plot
<code>title()</code>	Title (above plot)
<code>text()</code>	Text in plot
<code>mtext()</code>	Text in margin
<code>legend()</code>	List of symbols

These are all *added* to existing plots.

### Graphical parameters

<code>pch</code>	Symbol ( <i>plotting character</i> )
<code>mfrow, mfcol</code>	Several plots on one ( <i>multiframe</i> )
<code>xlim, ylim</code>	Plot limits
<code>lty, lwd</code>	Line type/width
<code>col</code>	Colour
<code>cex, mex</code>	Character size and line spacing in margins

See the help page for `par` for more details.

# Programming

Conditional execution	<pre>if(p&lt;0.05)   print("Hooray!")</pre>
— with alternative	<pre>if(p&lt;0.05)   print("Hooray!") else   print("Bah.")</pre>
Loop over list	<pre>for(i in 1:10)   print(i)</pre>
Loop	<pre>i &lt;- 1 while(i&lt;10) {   print(i)   i &lt;- i + 1 }</pre>
User-defined function	<pre>f &lt;- function(a,b,doit=FALSE){   if (doit)     a + b   else     0 }</pre>
<p>In flow control one uses <code>a &amp;&amp; b</code> and <code>a    b</code> where <code>b</code> is only computed if necessary; that is, if <code>a</code> then <code>b</code> else <code>FALSE</code> and if <code>a</code> then <code>TRUE</code> else <code>b</code></p>	