# Input and Output

#### Text

- A string of characters is said to be of mode character.
- Character strings are denoted by either double quotes " " or single quotes '.'
- Strings can be arranged into vectors and matrices just like numbers.
- We can paste string using paste(???, sep).

```
x <- "Citroen SM"
y <- "Jaguar XK150"
z <- "Ford Falcon GT-H0"
(wish.list <- paste(x, y, z, sep=", "))</pre>
```

- ## [1] "Citroen SM, Jaguar XK150, Ford Falcon GT-HO"
  - Special characters with the escape character .
  - \n for newline, \t for tab, \b for backspace, \" for ", \\ for \
  - If a character string can be understood as a number, the as.numeric(x) is used.

```
as.numeric("10.1")
```

#### ## [1] 10.1

• Use as.character(x) to coerce a number into a character string.

```
as.character(10.1)
```

```
## [1] "10.1"
```

## Input from a file

- R provides a number of ways to read data from a file.
- scan function is the most flexible one.
- scan to read a vector of values from a file.
- scan(file= "", what=0, n=-1, sep="", skip=0, quiet=FALSE)
- all parameters are optional
- file gives the file to read from.
  - The default " " indicates read from the keyboard.
- what gives an example of the mode of data to be read.
  - use 0 for numeric value, use " " for character data.
- n gives the number of elements to read.
  - if n = -1 then scan keeps reading until EOF.
- sep allows you to specify that is used to separate values such as ","
- skip is the number of lines to skip before start reading.
- quiet controls whether or not scan reports how may values it has read.

# Example: file input

```
# You need to change the path of the file accordingly.
data <- scan(file="data/data1.txt")
n <- length(data)
data.sort <- sort(data)
data.1qrt <- data.sort[ceiling(n/4)]
data.med <- data.sort[ceiling(n/2)]</pre>
```

#### Input from the keyboard

• scan can be used to read from the keyboard if file is "".

```
scan(file="", what="") # character input
scan(file="", what=0) # numeric input
```

- readline(prompt) read a single line of text from the keyboard.
- prompt (default ""): takes the optional character input

```
your_name <- readline("Input your name : ")</pre>
```

## Example: Root of quadratic

```
cat("find zeros of a2*x^2 + a1*x + a0 = 0\n")
a2 <- as.numeric(readline("a2= "))
a1 <- as.numeric(readline("a1= "))
a0 <- as.numeric(readline("a0= "))
discrim <- a1^2 - 4*a2*a0
if (discrim > 0) {
 roots \leftarrow (-a1 + c(1,-1) * sqrt(a1^2 - 4*a2*a0))/(2*a2)
} else if (discrim == 0) {
  roots <- -a1/(2*a2)
} else {
  roots <- c()
}
if (length(roots) == 0){
 cat("no root\n")
} else if (length(roots)==1){
  cat("single root at", roots, "\n")
  cat("roots at", roots[1], "and", roots[2], "\n")
```

# Output to a file

- Generally use write or write.table for writing numeric.
- cat for writing text, or a combination of numeric and text.
- write(x, file = "data", ncolumns = if(is.character(x)) 1 else 5, append = FALSE)
- x is the vector to be written.
  - If x is a matrix then it is converted to a vector (column by column)
- file gives the file to write as a character string
  - default "data" writes to a file called data in the current working directory
  - to write to the screen use file=""
- ncolumns: the number of columns to write the data in.
- append indicates whether to append to or overwrite the file.
- write converts matrices to vectors
- Since R stores its matrices by column, you should transpose the matrix to write if you want the output to reflect the matrix structure.

```
( x <- matrix(1:24, nrow=4, ncol=6))
write(t(x), file = "out.txt", ncolumn=6)</pre>
```

#### cat for writing to a file

- cat is more flexible command.
- cat(???, file="", sep="", append=FALSE)
- ??? is a list of expressions (separated by commas) that are coerced into character strings, concatenated, and then written.
- file gives the file to write or append to as a character string.
  - the default "" writes to the screen.
- sep is character string that is inserted between objects.
- append indicates whether to append to or overwrite the file.

# dump

• dump creates a text representation of almost any object that can subsequently read by source.

```
x <- matrix(rep(1:5, 1:5), nrow=3, ncol=5)
dump("x", file="result.txt")
rm(x)
source("result.txt")
x</pre>
```

```
## [,1] [,2] [,3] [,4] [,5]
## [1,] 1 3 4 4 5
## [2,] 2 3 4 5 5
## [3,] 2 3 4 5 5
```

#### Plotting

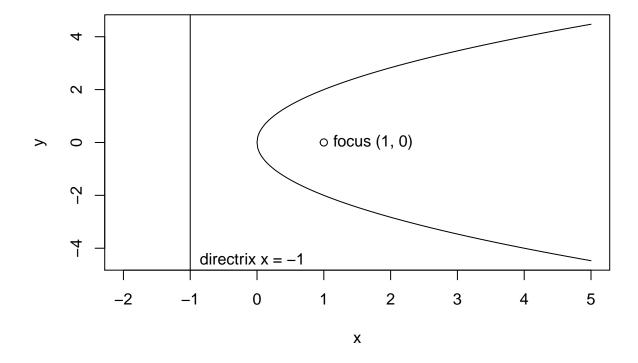
• We have already seen plot(x, y, type).

- To add points to the current plot, use points(x, y).
- To add lines, use lines(x, y).
- To add vertical or horizontal liens, use abline(v=xpos) or abline(y=ypos).
- points and lines take optional input col, which determines color.
- colors() show the complete list of colors.
- To add text labels[1] at (x[i], y[i]), use text(x, y, lables).
- pos is used to indicate the position. (See help(text))
- title(main) for the title (main is a character string).

# plotting a parabola $y^2 = 4x$

```
x <- seq(0, 5, by=0.01)
y.upper <- 2*sqrt(x)
y.lower <- -2*sqrt(x)
y.max <- max(y.upper)
y.min <- min(y.lower)
plot(c(-2,5), c(y.min, y.max), type="n", xlab="x", ylab="y")
lines(x, y.upper)
lines(x, y.lower)
abline(v=-1)
points(1,0)
text(1, 0, "focus (1, 0)", pos=4)
text(-1, y.min, "directrix x = -1", pos=4)
title("The parabola y^2 = 4*x")</pre>
```

# The parabola $y^2 = 4x$



# more than one plots

- windows() open additional graphics devices.
- par(mfrow = c(nr, nc)) or par(mfcol = c(nr, nc)) creates a grid of plots.

- mfrow = c(nr, nc)): with nr rows and nc columns, filled row by row.
- mfcol = c(nr, nc)) : filled column by column.

```
par(mfrow=c(2,2))
curve(x*sin(x), from=0, to=100, n=1001)
curve(x*sin(x), from=0, to=10, n=1001)
curve(x*sin(x), from=0, to=1, n=1001)
curve(x*sin(x), from=0, to=0.1, n=1001)
```

