```
rintro-commands.r
# first part of rintro.pdf...
# Basics
x <- 3
print(x)
          # NOTE: the rintro.pdf document uses "_" instead of "=".
# "=" is the correct symbol! (synonym for "<-")</pre>
y <- "Hello there"
y <- sqrt(10)
z <- x + y
           # use this to quit
# A basic R object is the "vector". R always prints some index hints
# to the left of a printout of a vector.
# A scalar in R is just a vector of length 1, so we still get the
# index hint to the left of the printout
x <- 1:5
print(x)
xx <- 1:50
x[1] <- 17
x[3:5] <- 0
w < -x[-3]
y \leftarrow c(1,5,2,4,7)
y[2]
```

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y[-3]
y[c(1,4,5)]
i <- 1:3
z \leftarrow c(9,10,11)
y[i] <- z
y <- y^2
y <- 1:10
y <- log(y)
y \leftarrow exp(y)
x <- 1:10
y <- c(5,4,3,2,1,5,4,3,2,1)
z <- x + y
x <- 1:10
y \leftarrow c(5,4,3,2,1,5,4,3,2,1)
x == 2
z < -(x == 2)
z <- (x<5); z
                     # ";" separates commands on the same line
                     # you can break an expression across lines
                     # if the expression is incomplete on the current line
x[x<5] \leftarrow y[x<5]
###########
# matrices are 2-dimensional arrays of elements of the same type
junk <- c(1,2,3,4,5, 0.5, 2, 6, 0, 1, 1, 0)
m <- matrix(junk,ncol=3)</pre>
m <- matrix(junk,ncol=3,byrow=T)</pre>
dim(m)
```

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y <- m[,1]
x < -m[,2]
z \leftarrow m[1,2]
zz <- t(m)
                  # matrix transpose!
ZZ
new <- matrix(1:9,ncol=3)</pre>
hello <- z + new
hello
m[1,3]
subm <- m[2:3,2:4]
m[1,]
m[2,3] < -7
m[2,3]
m[,c(2,3)]
m[-2,]
############
# Something slightly statistical to illustrate built-in functions
x \leftarrow runif(100,0,1) \# random sample of size 100 from Unif[0,1]
mean(x)
y <- mean(y)
help(mean)
min(x)
max(x)
summary(x)
help(summary)
                    # most commands in R have a help page!
############
# aside: a vector can only have elements of one type in it.
         a matrix can only have elements of one type in it.
```

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# use a **list**.
# if you want a matrix-like object with different data types in it,
# use a **data.frame** (which is basically a list of columns).
# data read into R from external files is usually stored in a
# data frame, by default.
# Here are som examples involving lists
who <- list(name="Joe", age=45, married=T)</pre>
who$name
who[[1]]
who$age
who[[2]]
who$married
who[[3]]
names(who)
who$name <- c("Joe", "Steve", "Mary")
who$age <- c(45,23)
who$married <- c(T,F,T)
#########
# there is more useful info, on loops and creating and using
# functions, in the rintro.pdf handout!
#########
```

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if you want a vector-like object with different data types in it,