- * A program is a set of instructions for a computer to follow
- Programs are often used to manipulate data (in all type and formats you discussed last week)
- Simple to complex
 - the scripts you wrote last week (simple)
 - instructions to analyze relationships in census data and visualize them
 - a model of global climate

- Programs can be written in many different languages
 (all have their strengths and weakness)
- Languages expect instructions in a particular form (syntax) and then translate them to be readable by the computer
- * Languages have evolved to make it help users write programs that are easy to understand, re-use, extend, test, run quickly, use lots of data...

- Operations (=,+,-,...concatenate, copy)
- * Data structures (simple variables, arrays, lists...)
- Control structures (if then, loops)
- * Modules...

Concepts common to all languages through the syntax may be different

Modularity

Main controls the overall flow of program- calls to the functions/ modules/building blocks



- * A program is often multiple pieces put together
- * These pieces or modules can be used multiple times

- Modularity
 - breaking your instructions down into individual pieces
 - identifying instructions that can be reused
 - an ecosystem model might re-use instructions for calculating how a species grows
 - an accounting program might re-use instructions for computing net present value from interest rates
 - modules often become 'black boxes' which hides detail that might make understanding the program overly complex
 - most languages have lots of black boxes already written and most allow you to write your own

Best practices for software development

- Read: Wilson G, Aruliah DA, Brown CT, Chue Hong NP, Davis M, et al. (2014) Best Practices for Scientific Computing. PLoS Biol 12(1): e1001745. doi:10.1371/journal.pbio.1001745
- Blanton, B and Lenhardt, C 2014. A Scientist's Perspective on Sustainable Scientific Software. Journal of Open Research Software 2(1):e17, DOI: http://dx.doi.org/10.5334/jors.ba
- but also
- * <u>http://simpleprogrammer.com/2013/02/17/principles-are-</u> <u>timeless-best-practices-are-fads/</u>

Box 1. Summary of Best Practices

- 1. Write programs for people, not computers.
- (a) A program should not require its readers to hold more than a handful of facts in memory at once.
- (b) Make names consistent, distinctive, and meaningful.
- (c) Make code style and formatting consistent.
- 2. Let the computer do the work.
- (a) Make the computer repeat tasks.
- (b) Save recent commands in a file for re-use.
- (c) Use a build tool to automate workflows.
- 3. Make incremental changes.
- (a) Work in small steps with frequent feedback and course correction.
- (b) Use a version control system.
- (c) Put everything that has been created manually in version control.
- 4. Don't repeat yourself (or others).
- (a) Every piece of data must have a single authoritative representation in the system.
- (b) Modularize code rather than copying and pasting.
- (c) Re-use code instead of rewriting it.

- 5. Plan for mistakes.
- (a) Add assertions to programs to check their operation
- (b) Use an off-the-shelf unit testing library.
- (c) Turn bugs into test cases.
- (d) Use a symbolic debugger.
- 6. Optimize software only after it works correctly.
- (a) Use a profiler to identify bottlenecks.
- (b) Write code in the highest-level language possible.
- 7. Document design and purpose, not mechanics.
- (a) Document interfaces and reasons, not implementation
- (b) Refactor code in preference to explaining how it we
- (c) Embed the documentation for a piece of software in software.
- 8. Collaborate.
- (a) Use pre-merge code reviews.
- (b) Use pair programming when bringing someone new speed and when tackling particularly tricky problem
- (c) Use an issue tracking tool.

Best practices for model (software) development

- Common problems
 - Unreadable code (hard to understand, easy to forget how it works, hard to find errors, hard to expand)
 - Overly complex, disorganized code (hard to find errors; hard to modify-expand)
 - Insufficient testing (both during development and after)
 - Not tracking code changes (multiple versions, which is correct?)

Steps for building model

- * We are going to use R; but the basic design of programs are similar across many programming languages
- * Why R?
 - * Free (and open source) software
 - * Good (and getting better) visualization tools
 - * Growing user community who make their R code available
 - (currently 2800+ user packages on CRAN R server)
 - * Links with other tools and languages (GIS, Python, C, C++...)
 - * Built in tools to deal with space and time
 - Lots of user support

Steps for building model

- * Why not R?
 - Not particularly computationally efficient (e.g slow for repetitive computations) ; hard to parallelize
 - Not the right tool for developing really complex models (you don't develop GCMs in R!)

STEPS: Program Design

- 1. Clearly define your goal as precisely as possible, what do you want your program to do
 - 1. inputs/parameters
 - 2. outputs
- 2. Implement and document
- 3. Test
- 4. Refine



- 1. Design the program "conceptually" "on paper" in words or figures
- 2. Translate into a step by step representation
- 3. Choose programming language
- 4. Define inputs (data type, units)
- 5. Define output (data type, units)
- 6. Define structure
- 7. Write program
- 8. Document the program
- 9. Test the program
- 10. Refine...

Best practices for model (software) development

- Let us change our traditional attitude to the construction of programs: Instead of imagining that our main task is to instruct a computer what to do, let us concentrate rather on explaining to humans what we want the computer to do. --Donald E. Knuth, Literate Programming, 1984
- Developing readable (by PEOPLE) code and documenting what you are doing is essential
- * "When was the last time you spent a pleasant evening in a comfortable chair, reading a good program?"— Bentley (1986)

Best practices for software development

- * Automated tools (useful for more complex code development
- * (note that GP's often create programs > 100 lines of code)
- Automated documentation
 - http://www.stack.nl/~dimitri/doxygen/
 - http://roxygen.org/roxygen2-manual.pdf
- Automated test case development
 - http://r-pkgs.had.co.nz/tests.html
- * Automated code evolution tracking (Version Control)
 - https://github.com/

Designing Programs

- * Inputs sometimes separated into input data and parameters
 - * input data = the "what" that is manipulated
 - parameters determine "how" the manipulation is done
 - * "sort -n file.txt"
 - * sort is the program set of instructions its a black box
 - * input is file.txt
 - parameters is -n
 - output is a sorted version of file.txt
 - * my iphone app for calculating car mileage
 - * inputs are gallons and odometer readings at each fill up
 - * graph of is miles/gallon over time
 - parameters control units (could be km/liter, output couple be presented as a graph or an average value)

Designing Programs

- What's in the box (the program itself) that gives you a relationship between outputs and inputs
 - * the link between inputs and output
 - breaks this down into bite-sized steps or calls to other boxes)
 - think of programs as made up building blocks
 - * the design of this set of sets should be easy to follow

Building Blocks

- Instructions inside the building blocks/box
 - Numeric data operators
 - * +,-,/,*,%*%
 - * Strings
 - substr, paste..
 - * Math
 - * sin, cos, exp, min, max...
 - these are themselves programs boxes
 - * R-reference card is useful!

Best practices for software development

- * Structured practices that ensures
 - clear, readable code
 - modularity (organized "independent" building blocks)
 - * testing as you go and after
 - code evolution is documented

Building Blocks

- * Functions (or objects or subroutines)!
- The basic building blocks
- Functions can be written in all languages; in many languages (object-oriented) like C++, Python, functions are also objects
- Functions are the "box" of the model the transfer function that takes inputs and returns outputs
- More complex models made up of multiple functions; and nested functions (functions that call/user other functions)

Functions in R

* Format for a basic function in R

#' documentation that describes inputs, outputs and what the function does
FUNCTION NAME = function(inputs, parameters) {
 body of the function (manipulation of inputs)
 return(values to return)

In R, inputs and parameters are treated the same; but it is useful to think about them separately in designing the model - collectively they are sometimes referred to as arguments

ALWAYS USE Meaningful names for your function, its parameters and variables calculated within the function

A simple program: Example

- * Input: Reservoir height and flow rate
- * Output: Instantaneous power generation (W/s)
- * Parameters: $K_{Efficiency}$, ϱ (density of water), g (acceleration due to gravity)

 $P = \varrho * h * r * g * K_{Efficiency}$

P is Power in watts, ϱ is the density of water (~1000 kg/m3), h is height in meters, r is flow rate in cubic meters per second, g is acceleration due to gravity of 9.8 m/s2, K_{Efficiency} is a coefficient of efficiency ranging from 0 to 1.



```
* Example (power_gen.R)
```

```
power_gen = function(height, flow, rho=1000, g=9.8, Keff=0.8) {
  result = rho * height * flow * g * Keff
  return(result)
}
```

Building Models

- * Inputs/parameters are height, flow, rho, g, and K
- * For some (particularly parameters) we provide default values by assigning them a value (e.g Keff = 0.8), but we can overwrite these
- * Body is the equations between { and }
- * return tells R what the output is

```
power_gen = function(height, flow, rho=1000, g=9.8, Keff=0.8) {
  result = rho * height * flow * g * Keff
  return(result)
}
```

Building Models: Using the model

```
> power_gen(20,1)
[1] 156800
> power_gen(height=20,flow=1)
[1] 156800
> power.guess = power_gen(height=20,flow=1)
> power.guess
[1] 156800
> power.guess = power_gen(flow=1, height=20)
> power.guess
[1] 156800
```

Arguments to the function follow the order they are listed in your definition Or you can specify which argument you are referring to when you call the program

```
power_gen = function(height, flow, rho=1000, g=9.8, K=0.8) {
# calculate power
result = rho * height * flow * g * K
return(result)
}
```

Building Models

- * Always write your function in a text editor and then copy into R
- * By convention we name files with functions in them by the name of the function.R
 - * so power_gen.R
- you can also have R read a text file by *source("power_gen.R")* make sure you are in the right working directory
- Eventually we will want our function to be part of a package (a library of many functions) - to create a package you must use this convention (name.R)

Building Models: Using the model

```
> power_gen(height=20, flow=1)
[1] 156800
> power_gen(height=20, flow=1, Keff=0.8)
[1] 156800
> power_gen(height=20, flow=1, Keff=0.5)
[1] 98000
> power_gen(height=10, flow=1, Keff=0.5)
[1] 49000
```

Defaults take the value they were assigned in the definition, but can be overwritten

```
power_gen = function(height, flow, rho=1000, g=9.8, K=0.8) {
# calculate power
result = rho * height * flow * g * K
return(result)
```

}



The scope of a variable in a program defines where it can be "seen"

Variables defined inside a function cannot be "seen" outside of that function

There are advantages to this - the interior of the building block does not 'interfere' with other parts of the program

```
> power_gen
function(height, flow, rho=1000, g=9.8, K=0.8) {
```

```
# calculate power
result = rho * height * flow * g * K
return(result)
}
> result
Error: object 'result' not found
> K
Error: object 'K' not found
```

>

One of the equations used to compute automobile fuel efficiency is as follows this is the power required to keep a car moving at a given speed

```
Pb = c_{rolling} * m *g*V + 1/2 A*p_{air}*c_{drag}*V^3
```

where $c_{rolling}$ and c_{drag} are rolling and aerodynamic resistive coefficients, typical values are 0.015 and 0.3, respectively. V: is vehicle speed (assuming no headwind) in m/s (or mps) m: is vehicle mass in kg A is surface area of car (m2) g: is acceleration due to gravity (9.8 m/s2) p_{air} = density of air (1.2kg/m3) Pb is power in Watts

Write a function to compute power, given a truck of m=31752 kg (parameters for a heavy truck) for a range of different highway speeds plot power as a function of speed how does the curve change for a lighter vehicle

Note that 1mph=0.477m/s

Simple Functions

Note that we can use vectors (list of numbers) in addition to single numbers as inputs - see use of "v"

lines(v, power(V=0.447*v, m=61752, A=25))

```
power = function(cdrag=0.3, crolling=0.015,pair=1.2,g=9.8,V,m,A) {
P = crolling*m*g*V + 1/2*A*pair*cdrag*V**3
return(P)
}
v=seq(from=0, to=100, by=10)
plot(v, power(V=0.447*v, m=31752, A=25))
```

Simple Functions

```
Power Required by Speed
#'
#'
  This function determines the power required to keep a vehicle moving
#'
at
#' a given speed
  @param cdrag coefficient due to drag default=0.3
#'
#' @param crolling coefficient due to rolling/friction default=0.015
#' @param v vehicle speed (m/2)
#' @param m vehicle mass (kg)
#' @param A area of front of vehicle (m2)
#' @param g acceleration due to gravity (m/s) default=9.8
#' @param pair (kg/m3) default =1.2
#' @return power (W)
power = function(cdrag=0.3, crolling=0.015,pair=1.2,g=9.8,V,m,A) {
P = crolling*m*g*V + 1/2*A*pair*cdrag*V**3
return(P)
}
v=seq(from=0, to=100, by=10)
plot(v, power(V=0.447*v, m=31752, A=25))
lines(v, power(V=0.447*v, m=61752, A=25))
```

- Understanding data types is important for designing your model I/O; specifying what the model will do
- Data types and data structures are necessary for creating more complex inputs and outputs
- * All programming languages have sets of data types
 - single values: character, integer, real, logical/boolean (Y/N)
 - data structures: arrays, vectors, matrices,
 - * in R core types; dataframes, lists, factors
 - * in R defined types: spatial, date...



Building Programs

A core issue in modeling (both designing and using) are the **data structures** / formats used to hold data that is input and output from programs: In good programs, data structures support organization and program flow and readability Key Programming concepts: Data types and structures

- Good data structures are:
 - * as simple as possible
 - easy understand (readable names, and sub-names)
 - easy to manipulate (matrix operations, applying operations by category)
 - easy to visualize (graphs and other display)

- Vectors a 1-dimensional set of numbers
- * a = c(1,5,8,4,22,33)
- Matrix a 2-dimensional set of numbers (organized in rows and columns)
- * b = matrix(a, nrow=2, ncol=3)

```
> a = c(1,5,8, 4, 22,33)
>
> b = matrix(a, nrow=2, ncol=3)
> a
[1] 1 5 8 4 22 33
> b
      [,1] [,2] [,3]
[1,] 1 8 22
[2,] 5 4 33
```

- You can also define an "empty" matrix to fill values in later
- think of creating a data structure to store energy production in winter and summer for 6 different power plants)
- * res = matrix(nrow=2, ncol=6)

```
> res = matrix(nrow=2, ncol=6)
> res
     [,1] [,2] [,3] [,4] [,5] [,6]
      NA
           NA
                 NA
                      NA
                           NA
[1,]
                                NA
[2,]
      NA
           NA
                 NA
                      NA
                           NA
                                NA
>
```

- * You can combine vectors into a matrix using
 - * cbind by columns
 - * *rbind* by rows

- * A really useful data structure in R is a data frame
- Dataframe's are like matrices = they have rows and columns but they don't have to be numeric (although they can be)
- * Useful if you have data that is of mixed type

Data Frame Creation Example

We often want to "create" data to explore ideas/function behavior

```
>
> mth.names = c("Jan", "Feb", "Mar", "Apr", "May",
+ "Jun","Jul","Aug","Sep","Oct","Nov","Dec")
>
> reservoir.operation = data.frame(month=mth.names)
>
> reservoir.operation
   month
     Jan
1
2
     Feb
3
     Mar
4
     Apr
5
     May
6
     Jun
7
     Jul
8
     Aug
9
     Sep
10
     0ct
11
     Nov
12
     Dec
~
```

Data Frame Creation Example

```
>
>
> reservoir.operation$height = c(seq(from=22,to=10, by=-2), seq(from=12,to=20,
by=2))
> \#reservoir.operationheight = c(20, 18, 16, 14, 12, 10, 12, 14, 16, 18, 20)
>
> reservoir.operation$flowrate = rnorm(n=12, mean=3, sd=0.25)
>
> reservoir.operation
   month height flowrate
                                             Adding columns
     Jan
            22 2.967183
1
2
    Feb 20 2.923782
3
    Mar 18 2.891444
4
    Apr
            16 3.048090
                           seq - a sequence of number from to by
5
    May
            14 2.598533
6
    Jun
            12 3.027024
                           rnorm - generate, n numbers from a normal distribution
7
            10 2.906364
    Jul
                           with a given mean and standard deviation
8
    Aug
            12 3.460389
             14 2.965138
9
     Sep
10
    0ct
             16 3.320663
11
    Nov
            18 3.300340
12
             20 2.916503
    Dec
>
```

- Of course we can use matrices/data frames as inputs/ output for our models
- Example using our power_gen model from earlier using vectors instead of single values

```
> > power_gen(height=reservoir.operation$height, flow=reservoir.operation$flow)
[1] 511779.6 458449.1 408040.5 382352.4 285215.0 284782.4 227858.9 325553.4
[9] 325453.5 416544.0 465744.0 457307.7
> power_gen
function (height, flow, rho = 1000, g = 9.8, Keff = 0.8)
{
    result = rho * height * flow * g * Keff
    return(result)
}
```

- * Why does this work?
- Because height, flow columns are both from reservoir.operation (a data frame) so they are vectors of the SAME length
- * So when you multiply height* flow, you multiply
 - * height[1]*flow[1],,, and then height[2]*flow[2] etc

```
> > power_gen(height=reservoir.operation$height, flow=reservoir.operation$flow)
[1] 511779.6 458449.1 408040.5 382352.4 285215.0 284782.4 227858.9 325553.4
[9] 325453.5 416544.0 465744.0 457307.7
> power_gen
function (height, flow, rho = 1000, g = 9.8, Keff = 0.8)
{
    result = rho * height * flow * g * Keff
    return(result)
}
```

- * Matrix multiplication is different
- * in R, this would be
 - * k %*% m



* Matrix multiplication is often used within certain types of models...we will get to examples later

Data Frame Creation Example

```
>
>
> reservoir.operation$height = c(seq(from=22,to=10, by=-2), seq(from=12,to=20,
by=2))
> \#reservoir.operationheight = c(20, 18, 16, 14, 12, 10, 12, 14, 16, 18, 20)
>
> reservoir.operation$flowrate = rnorm(n=12, mean=3, sd=0.25)
>
> reservoir.operation
   month height flowrate
                                             Adding columns
            22 2.967183
     Jan
1
2
    Feb 20 2.923782
3
    Mar 18 2.891444
4
    Apr
            16 3.048090
5
    May
            14 2.598533
6
    Jun
            12 3.027024
7
            10 2.906364
    Jul
8
    Aug
            12 3.460389
             14 2.965138
9
     Sep
    0ct
             16 3.320663
10
11
    Nov
            18 3.300340
12
             20 2.916503
     Dec
>
```

* We can also use data frames (or matrices) to store results

> power_gen(height=reservoir.operation\$height, flow=reservoir.operation\$flow)
[1] 511779.6 458449.1 408040.5 382352.4 285215.0 284782.4 227858.9 325553.4
[9] 325453.5 416544.0 465744.0 457307.7

> reservoir.operation\$power = power_gen(height=reservoir.operation\$height,
flow=reservoir.operation\$flow)

> reservoir.operation

	month	height	flowrate	power
1	Jan	22	2.967183	511779.6
2	Feb	20	2.923782	458449.1
3	Mar	18	2.891444	408040.5
4	Apr	16	3.048090	382352.4
5	May	14	2.598533	285215.0
6	Jun	12	3.027024	284782.4
7	Jul	10	2.906364	227858.9
8	Aug	12	3.460389	325553.4
9	Sep	14	2.965138	325453.5
10	0ct	16	3.320663	416544.0
11	Nov	18	3.300340	465744.0
12	Dec	20	2.916503	457307.7

- Some other useful commands
 - *with* allows you to use the names of columns in the data frame directly
 - *summary* summaries of columns (max, min, mean...)

> with(reservoir.operation, barplot(power, names=month, ylab="Power (W/s)"))
>



Summary

>

>

>

>

> summary(reservoir.operation)

month		height	flowrate	power
Apr	:1	Min. :10.0	Min. :2.599	Min. :227859
Aug	:1	1st Qu.:13.5	1st Qu.:2.914	1st Qu.:315394
Dec	:1	Median :16.0	Median :2.966	Median :395196
Feb	:1	Mean :16.0	Mean :3.027	Mean :379090
Jan	:1	3rd Qu.:18.5	3rd Qu.:3.111	3rd Qu.:457593
Jul	:1	Max. :22.0	Max. :3.460	Max. :511780
(Other):6				

We can also use other functions and built in R functions
 (like *mean*, *lm*, *sum*) within our function

You can imbed any function you write (or already in R) in your function

Key Programming concepts: functions calling other functions

* Example of embedded function

>

* Lets say we want to commute total annual power generated, given our inputs of average height and flow for each month?

```
* what additional information would we need?
```

```
>
> power_gen(height=reservoir.operation$height, flow=reservoir.operation$flow)
[1] 511779.6 458449.1 408040.5 382352.4 285215.0 284782.4 227858.9 325553.4
[9] 325453.5 416544.0 465744.0 457307.7
> power_gen
function (height, flow, rho = 1000, g = 9.8, Keff = 0.8)
{
    result = rho * height * flow * g * Keff
    return(result)
}
```

```
#' Total Power Generation
#'
#' This function computes total power generation from a reservoir given its
height and flow rate into turbines and number of days (and secs) within
those days that the turbines are in operation
#' @param rho Density of water (kg/m3) Default is 1000
#' @param g Acceleration due to gravity (m/sec2) Default is 9.8
#' @param Keff Turbine Efficiency (0-1) Default is 0.8
#' @param height height of water in reservoir (m)
#' @param flow flow rate (m3/sec)
#' @param number of days
#' @param secs in days Default is 86400
#' @author Naomi
#' @examples power_gen(20, 1, 10)
#' @return Power generation (MW)
power gen total = function(height, flow, days, secs=86400, rho=1000, g=9.8,
Keff=0.8) {
result = rho * height * flow * g * Keff
result = result * days * secs
total = sum(result)/1e6
return(total)
}
> power_gen_total(reservoir.operation$height, reservoir.operation$flowrate,
days=30)
[1] 11702915
>
```

Data Structures

- * vector, (c)
- * matrices, arrays
- * data frames
- * lists
- * factors

- * Lists are the most "informal" data structures in R
- * List are really useful for keeping track of and organizing groups of things that are not all the same
- * A list could be a table where number of rows is different for each column
- A list can have numeric, character, factors all mixed together
- List are often used for returning more complex information from function (e.g. lm)

* A simple list: using names to identify elements

```
> sale = list(number=2, quality="high", what="apple", cost=4)
> sale
$number
[1] 2
$quality
[1] "high"
```

```
$what
[1] "apple"
```

\$cost [1] 4

```
> costs = c(20,40,22, 32, 5)
> quality = c("G", "G", "F", "G", "B")
> purchased = c(33,5,22,6,7)
> sales = data.frame(costs=costs, quality=quality, purchased=purchased)
> sales
                     >costs = c(73,44)
costs quality purchased
                     >quality = c("G","G")
1
  20
        G
            33
                      >purchased = c(100, 22)
2
     G 5
 40
3
 22 F 22
                      >sales2 = data.frame(costs=costs, quality=quality
                      purchased=purchased)
 32 G 6
4
5
   5
       В
            7
```

With lists we can combine sales data frames from two different places into a single data structure

Lists

```
>
> markets = list(site1=sales, site2=sales2)
> markets
$site1
  costs quality purchased
     20
1
                         33
               G
2
     40
               G
                          5
3
4
               F
   22
                         22
                          6
     32
               G
5
                          7
      5
               В
$site2
  costs quality purchased
1
     73
                        100
               G
2
                         22
     44
               G
> markets[[1]]$costs
[1] 20 40 22 32 5
>
> markets$site1$costs
[1] 20 40 22 32 5
>
```

Lists

>					
>					
>	market	ts[[1]]			
	costs	quality	purchased		
1	20	G	. 33		
2	40	G	5		
3	22	F	22		
4	32	G	6		
5	5	В	7		
>					
>	<pre>> markets[[2]]</pre>				
	costs	quality	purchased		
1	73	G	100		
2	44	G	22		
>					
<pre>> markets[[1]][1,3]</pre>					
[1] 33					
>					

[[]] is used to get elements from the list

 one of the most useful things to do with list is to use them to return multiple 'items' from a function

```
#' computes profit from price for forest plot and Mg/C in that plot
#' @param price ($)
#' @param carbon (MgC)
#' @return list with mean, min, and max prices
compute_carbonvalue = function(price, carbon) {
  cost.per.carbon = price/carbon
  a = mean(cost.per.carbon)
  b = max(cost.per.carbon)
  c = min(cost.per.carbon)
  result = list(avg=a, min=c, max=b)
  return(result)
}
```

* example: returning lists from a function

```
>
> obs = data.frame(prices=c(23,44,60,4,2,33,59),
forestC=c(59,88,100,10,8,79,300))
> obs
  prices forestC
      23
              59
1
2
              88
      44
3
      60
             100
4
       4
              10
5
       2
               8
6
      33
              79
7
      59
             300
> forest.res = compute_carbonvalue(obs$prices, obs$forestC)
> forest.res
$avg
[1] 0.3934598
$min
[1] 0.1966667
$max
[1] 0.6
```

>

example: returning lists from a function

```
> obs=data.frame(prices=c(18,2,12,5), grassC=c(22,3,19,8))
```

```
> grass.res=compute_carbonvalue(obs$prices, obs$grassC)
```

> grass.res
\$avg
[1] 0.6853569

\$min [1] 0.625

>

```
$max
[1] 0.8181818
```

- * Many functions that you use in R, return lists
- *names* (to see what is in a list)
- * *attributes* (to see what is in a list)

```
> names(forest.res)
[1] "avg" "min" "max"
> attributes(forest.res)
$names
[1] "avg" "min" "max"
```

lm is an
 example of a function that
 returns a list >> res = lm(obs\$prices~obs\$forestC) > names(res) [1] "coefficients" "residuals" "effects" [4] "rank" "fitted.values" "assign" [7] "qr" "df.residual" "xlevels" [10] "call" "terms" "model" > res\$coefficients (Intercept) obs\$forestC 14.9789368 0.1865644 > res\$model obs\$prices obs\$forestC 23 59 1 2 44 88 3 60 100 4 10 4 5 2 8 33 79 6 7 59 300

>

Data Structures

- * vectors (c)
- * matrices, arrays
- * data frames
- * lists
- * factors

- * Factors (a bit tricky, basically a vector of "things" that has different levels (classes); not really numeric - so you can't average them!)
- * But can be useful for doing "calculations" with categories

```
>
> a = c(1,5,2.5,9,5,2.5)
> a
[1] 1.0 5.0 2.5 9.0 5.0 2.5
> mean(a)
[1] 4.166667
> a = as.factor(c(1,5,2.5,9,5,2.5))
> mean(a)
[1] NA
Warning message:
In mean.default(a) : argument is not numeric or logical: returning NA
> a
[1] 1 5 2.5 9 5 2.5
Levels: 1 2.5 5 9
> summary(a)
  12.5 5
             9
   2
         2
             1
  1
```

summary can be used with factors to get frequencies in each category (or "level")

```
>
>
> species.recorded = c("butterfly","butterfly","mosquito","butterfly","
ladybug", "ladybug", "mosquito")
> species.recorded = as.factor(species.recorded)
> species.recorded
[1] butterfly butterfly mosquito butterfly ladybug ladybug
                                                                mosquit
0
Levels: butterfly ladybug mosquito
> summary(species.recorded)
butterfly ladybug mosquito
        3
                  2
                            2
> plot(species.recorded)
>
```



```
>
                                              > species.recorded = c("butterfly","butterfly","mosquito","butterfly","
                                              ladybug", "ladybug", "mosquito")
                                              > species.recorded = as.factor(species.recorded)
                                              > species.recorded
                                              [1] butterfly butterfly mosquito butterfly ladybug
                                                                                               ladybug
                                                                                                        mosquit
                                              Levels: butterfly ladybug mosquito
                                              > summary(species.recorded)
> mean(summary(species.recorded))
                                              butterfly ladybug mosquito
                                                      3
                                                               2
                                                                        2
[1] 2.333333
                                              > plot(species.recorded)
> max(summary(species.recorded))
                                              >
[1] 3
> sum(summary(species.recorded))
[1] 7
                                                                            You can "do things" (apply
> sum(species.recorded)
Error in Summary.factor(c(1L, 1L, 3L, 1L, 2L, 2L, 3L), na.rm = FALSE) :
                                                                             functions) to the summary
  sum not meaningful for factors
                                                                             (frequency of each "factor"
> species.recorded
[1] butterfly butterfly mosquito butterfly ladybug
                                                                                          level
                                                        ladybug
[7] mosquito
Levels: butterfly ladybug mosquito
> summary(species.recorded)[1]
butterfly
> summary(species.recorded)[2]
ladybug
      2
> summary(species.recorded)[3]
mosquito
       2
>
```

- * A simple model that takes advantage of factors
- A model to compute an index of species diversity from a list of recorded species

$$D = \Sigma (n / N)^2$$

where n is the number of individuals in each species, and N is total number

```
# '
   Simpson's Species Diversity Index
# '
# '
   Compute a species diversity index
# '
   @param species list of species (names, or code)
# '
   @return value of Species Diversity Index
# '
   @examples
   compute simpson index(c("butterfly","butterfly","mosquito","butterfly",
#' "ladybug", "ladybug")))
#' @references
  http://www.tiem.utk.edu/~gross/bioed/bealsmodules/simpsonDI.html
# '
```

compute_simpson_index = function(species) {

```
species = as.factor(species)
tmp = (summary(species)/sum(summary(species))) ** 2
diversity = sum(tmp)
return(diversity)
}
```

Data Structures

- a bit more on factors; a list of numbers can also be a factor but then they are not treated as actual numbers you could think of them as "codes" or addresses or..
- use *as.numeric* or *as.character* to go back to a regular
 vector from a factor

```
> items = c(1,5,1,5,6,3)
> mean(items)
[1] 3.5
> items = as.factor(c(1,5,1,5,6,3))
> mean(items)
[1] NA
Warning message:
In mean.default(items) : argument is not numeric or logical: returning
NA
> summary(items)
1356
2121
> tmp = as.numeric(items)
> tmp
[1] 1 3 1 3 4 2
> mean(tmp)
[1] 2.333333
>
```

* example: returning lists from a function

```
#' Describe diversity based on a list of species
#'
#' Compute a species diversity index
#' @param species list of species (names, or code)
#' @return list with the following items
#' \describe{
#' \item{num}{ Number of distinct species}
#' \item{simpson}{Value of simpson diversity index}
#' \item{dominant}{Name of the most frequently occuring species}
#'}
#'
  @examples
#'
compute_diversity(c("butterfly","butterfly","mosquito","butterfly","ladybug",
"ladybug")))
#' @references
#' http://www.tiem.utk.edu/~gross/bioed/bealsmodules/simpsonDI.html
compute_diversity = function(species) {
species = as.factor(species)
tmp = (summary(species)/sum(summary(species))) ** 2
diversity = sum(tmp)
nspecies = length(summary(species))
tmp = which.max(summary(species))
dominant = names(summary(species)[tmp])
return(list(num=nspecies, simpson=diversity, dominant=dominant))
}
```

Assignment

In your group, write a function that performs some data analysis that is likely to be useful for your project...

Enter this function in your organization's github space Read/create some data to test your function Also include the data in github Submit the link the repository