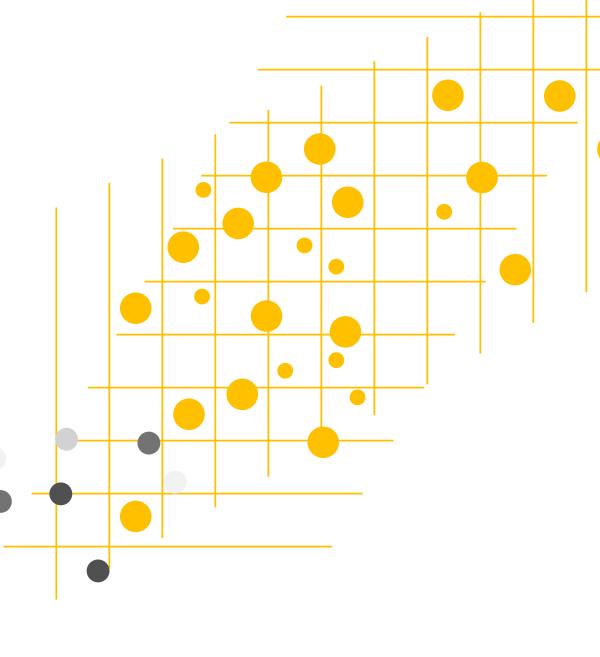
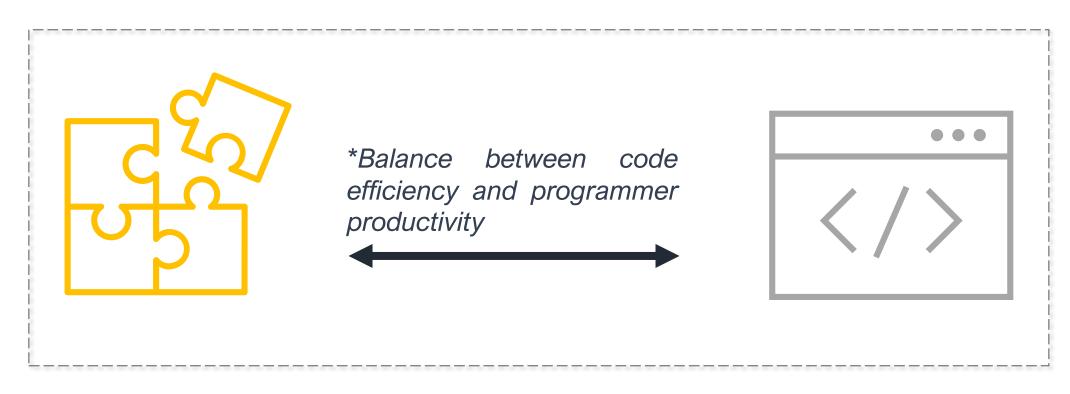
Measuring and Improving Performance

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What is performance?

How quickly can the computer undertake a particular task given a code



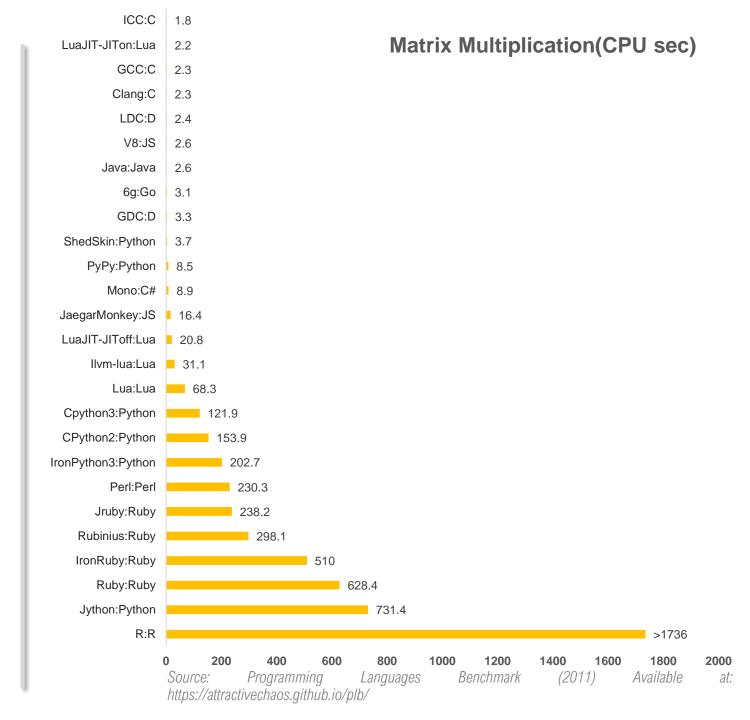
Is R an efficient coding language?



Language designed to make data analysis and statistics easier for people.

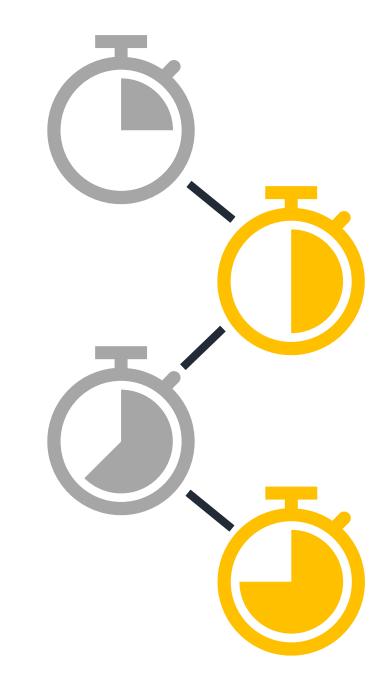


It is not a language designed with an efficient performance in mind.

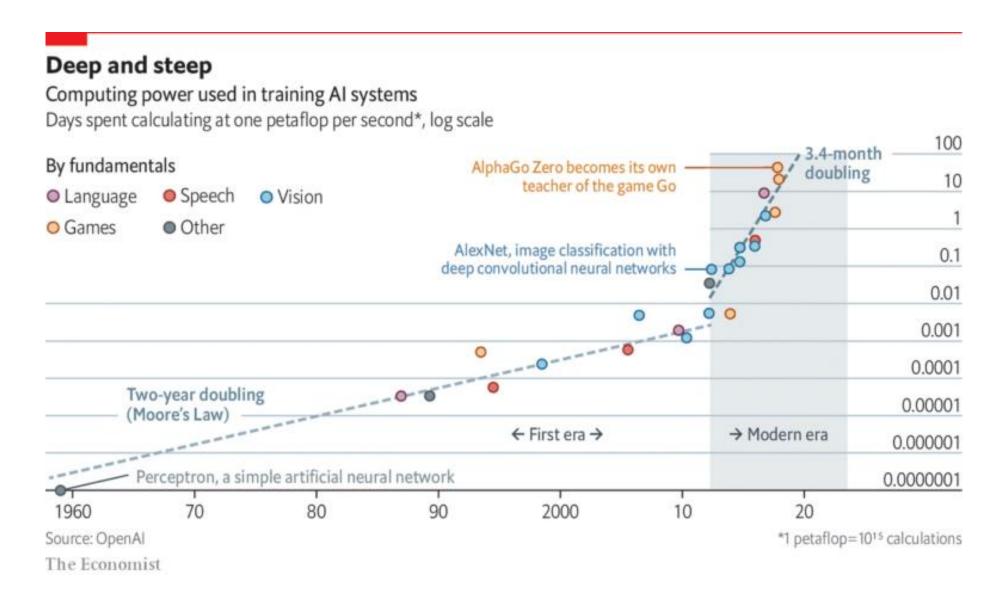


Why is improving performance important?

- The less efficient the code, the more time it takes processing data
 - i.e., reduce bottlenecks in your code
- Less memory usage
- CO2 emissions from energy usage with high intensity computing



Environmental Considerations



Environmental Considerations 👯 🗱 🗱











models is **Training** much more energy intensive than using them

"Aligning intelligence with climate mitigation" from Professor Lynn Kaack

artificial Relevant at intense Unlikely to affect your levels of computing, locally produced code as Al and machine change think Facebook and for class Google

Increasingly important learning continue to increase prevalence and usage

Google's machine translation system may process more than 100 billion words per day Facebook's datacenters are retrained anywhere from hourly to multimonthly

Important to keep in mind for your future career



How to make your code more efficient?

- 1. Start by identifying "bottle necks"
- 2. Experiment with alternatives to find faster code





1. Profiling

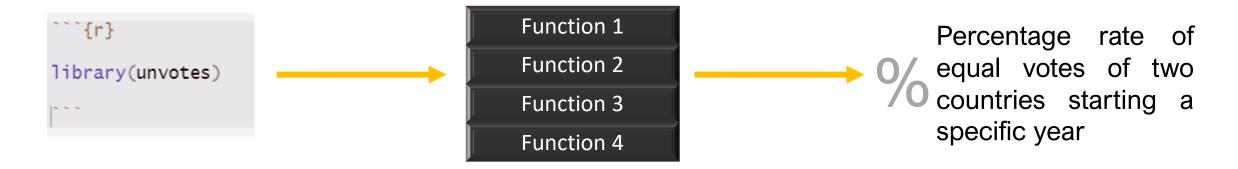
2. Benchmark

```
library(profvis)
library(bench)
```

Profiles

Your code profile will measure key factors in each line of code

- Run-time
- Memory
- Garbage collector



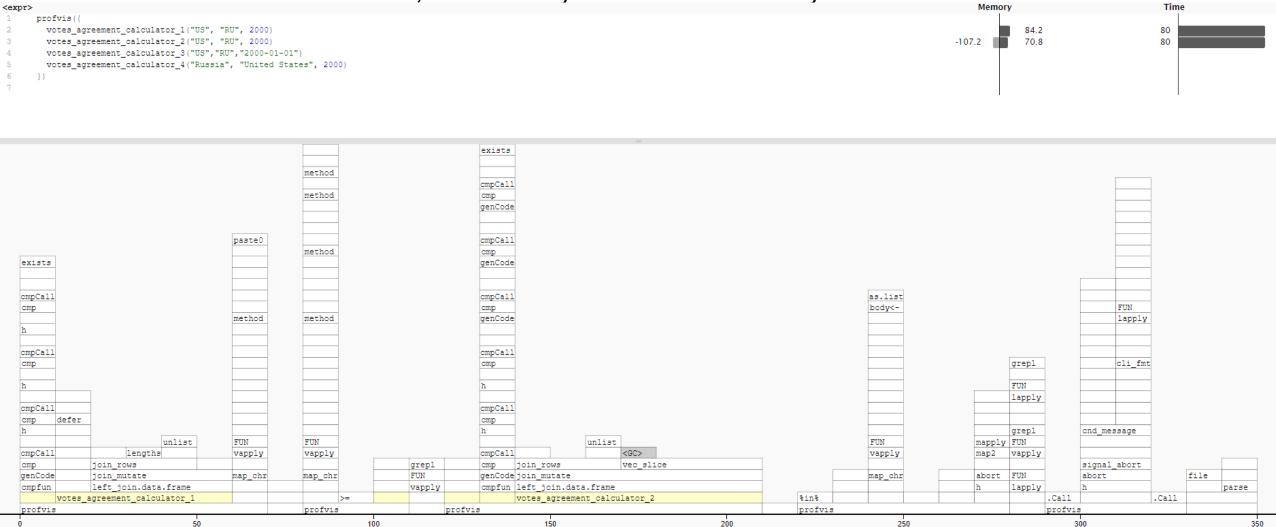
```
fr, message=FALSE, warning=FALSE}

profvis({
   votes_agreement_calculator_1("US", "RU", 2000)
   votes_agreement_calculator_2("US", "RU", 2000)
   votes_agreement_calculator_3("US", "RU", "2000-01-01")
   votes_agreement_calculator_4("Russia", "United States", 2000)
})
```

Visualizing Profiles

Vis output shows the source code, overlaid with bar graphs for memory and execution time for each line of code.

The bottom pane displays a flame graph showing the full call stack. It allows to see when objects are called more than one time. This is also displayed in the data tab, which let's you zoom interactively.



Benchmark

```
```{r, message=FALSE,warning=FALSE}
library(bench)
```

Always uses the highest precision APIs available for each operating systems (often nanoseconds)

Tracks memory allocations for each expression

Tracks the number and type of R garbage collections per expression iteration

Verifies equality of expression results by default, to avoid accidentally benchmarking inequivalent code

Uses adaptive stopping
by default, running
each expression for a
set amount of time
rather than for a
specific number of
iterations

Expressions are run in batches and summary statistics are calculated after filtering out iterations with garbage collections, to isolate the effects of garbage collection on running time.

The time and memory usage are returned as custom objects which have human readable formatting for display and comparisons.

There is also support for plotting with ggplot2, including custom scales.

#### Benchmark output

returns the results as a tibble

min, mean, median, mex, and itr/sec. These summarise the time taken by the expression. Focus on the minimum (best possible running time) and the median (the typical time).

Pay attention to the units. It is useful to know how many times a function needs to run before it takes a second. If a microbenchmark takes:

- \* 1 ms, then one thousand calls take a second
- \* 1 micros, then one million calls take a second
- \* 1 nanos, then one billion calls take a second

mem\_alloc tells you the amount of memory allocated by the first run, and n\_gc() tells you the total number of garbage collections over all runs. These are useful for assessing the memory usage of the expression.

#### What's next? **Improving** Performance

Approaches to improve

code

Code organization

Checking for existing solutions

Functions do as little work as possible

Vectorize code



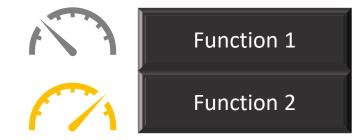
Balanced approach to code optimization

#### **Code Organization**

```
{r, message=FALSE,warning=FALSE}
library(bench)
```

#### Bench" package

Compare function speeds
 Which executes more quickly? How much memory is utilized?



- Example:
  - MDS Students' functions for Problem Set #2 for IDS

```
#Packages specifically for the example functions to work properly
library(tidyverse)
library(knitr)
library(dplyr)
library(countrycode)
library(tidyr)
library(lubridate)
library(rmarkdown)
library(unvotes)
```

#### Code Organization – Function #1

```
#Function #1
Create votes agreement calculator function
function 1 <- function(ccode a, ccode b, year min) {
 # Create joined data set with necessary columns; create year column from date column; select relevant v
 un <- un votes %>%
 left join(un roll calls, by = "rcid") %>%
 mutate(year = format(date, "%Y"), date = NULL) %>%
 select(rcid, country code, vote, year) %>%
 filter(!is.na(country code)) %>%
 filter(country code %in% c(ccode a, ccode b))
 # Pivot countries as columns and take values from the respective voting decision
 un votes wide <- un %>%
 pivot wider (names from = "country code", values from = "vote")
 # Calculate agreement share between ccode a and ccode b b since year min
 agreement share <- un votes wide %>%
 mutate(agreement = un votes wide[ccode a] == un votes wide[ccode b]) %>% #add agreement column
 filter(year >= year min, !is.na(agreement)) %>% #discard NAs in agreement variable
 summarise (agreement share = mean (agreement)) %>% #calculate mean() of agreement variable
 as.numeric() #convert to numeric
 return (round (agreement share, digits = 3))
```

#### Code Organization – Function #2

```
#Function #2
function 2 <- function(country 1, country 2, year min) {
 #Here we pull in the necessary data frames directly from the 'unvotes' package, assuming that the above
 un joined func <- un votes %>%
 inner join(un roll calls, by = "rcid") %>%
 group by (year = year (date), country)
 #Here we create a data frame specifically for country 1 and rename their "vote" column accordingly
 df 1 <- un joined func %>%
 filter(country %in% country 1, year >= year min) %>%
 rename(country 1 vote = vote)
 #We create a second data frame here for country 2 that does the same as above
 df 2 <- un joined func %>%
 filter(country %in% country 2, year >= year min) %>%
 rename (country 2 vote = vote)
 #Now, we merge these two new data frames into one so that we can compare their newly named vote columns
 joined dfs <- df 1 %>%
 inner join(df 2, by = "rcid")
 #Below, we count the number of times that the two countries agree with each other in the UN
 country vote agreement <- count(joined dfs, country_1_vote == country_2_vote)
 #Now we take the average and round it down to three decimal places
 average country vote agreement <- round((country vote agreement$n[2]/(country vote agreement$n[1]+cou
 #Lastly, we create a list including the two countries' names and how often they voted with each other
 returned list <- c(country 1, country 2, average country vote agreement)
 return (average country vote agreement)
```

#### Code Organization – Function Comparison

• Utilize "bench" to compare the two functions:

```
bench::mark(
 function_1("US", "RU", 2000),
 function_2("Russia", "United States", 2000)
)
```

Observe the results:

## Checking for Existing Solutions

- Outside resources:
  - Online resources:
    - Rseek at rseek.org
    - Stackoverflow at stackoverflow.com
    - CRAN Task Views at cran. Rstudio.com/web/views/
  - In-person resources:
    - Colleagues
    - Professors
    - Students

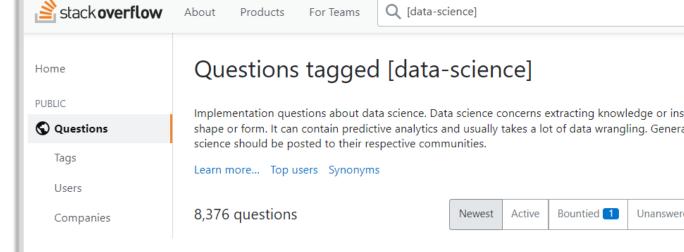


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About R R Homepage CRAN task views aim to provide some guidance which packages on CRAN are can also be automatically installed using the <a href="ctv">ctv</a> package. The views are intended and they are not meant to endorse the "best" packages for a given task.

To automatically install the views, the <u>ctv</u> package needs to be installed, e.g., visinstall.packages("ctv")

and then the views can be installed via install.views or update.views (where the ctv::install.views("Econometrics")

ctv::update.views("Econometrics")

To query information about a particular task view on CRAN from within R or to ctv::ctv("Econometrics")

ctv::available.views()

#### **Efficient Functions**



Have your functions do as little work as possible

Low runtime
Low memory usage



Some functions are faster than others







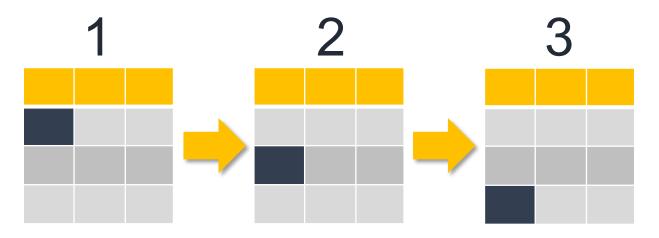
Requires experience and development of intuition

#### Examples:

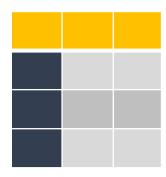
- \* Readr::read\_csv() is significantly faster than read.csv()
- \* rowSums(), colSums(), rowMeans(), and colMeans() are faster than apply() because they use vectors

#### Vectorize Your Code

For-loops iterate over a vector one at a time, increasing time and memory usage



Vectorized functions are applied to the entire vector at once rather than one at a time



Use "Microbenchmark" to compare simple functions like the following example

#### Vectorize Your Code - Example

```
#Function (vectorized) that multiplies two vectors.
vectorized vector <- function(n) {
 x \leftarrow 1:n
 v \leftarrow x * x
 return(v)
#Function (non-vectorized) that multiplies two vectors
vector for loop <- function(n) {
 x < -1:n
 v <- vector("numeric", length = n)</pre>
 for (i in 1:n) {
 y[i] \leftarrow x[i] * x[i]
 return(v)
#Here we use the microbenchmark function to show how many nanoseconds are needed to run each function 100
microbenchmark(vectorized vector(100),
 vector for loop(100),
 times = 100)
```

#### Vectorize Your Code – Example Output

- Observe how many nanoseconds are needed for each function
- Vectorized function is much more efficient and quicker

```
Unit: nanoseconds
expr min lq mean median uq max neval
vectorized_vector(100) 702 901 1227.05 1002.0 1301.0 8400 100
vector_for_loop(100) 8401 8701 9104.88 8801.5 9001.5 23301 100
```

#### Thanks!





