- $\hfill \ensuremath{\,^\circ}$ The function <code>system.time()</code> will do this for us.
- o R can directly save compressed.gz files with the file=gzfile("FileName.gz") option.
 - Using a csv.gz file could be faster or slower (depending on your storage subsystem speeds) but could save a lot of storage space.
- Let's save the data frame to temp2.csv.gz
 - The new file should be much smaller.
 - Let's read the .gz file to see if this worked.
 - R will automatically recognize the .gz extensions and decompress when loading.

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Using gzipped flat files is a great and simple way to transfer, load, and work with moderately large data sets.

- If speed or memory are an issue for your project when using R built-in I/O functions, you might need an external package.
- For example, there are many packages that can speed up the reading and writing speeds in R.

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 The vroom package comes with functions called vroom and vroom_write which are very efficient and speedy.

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- $\,\circ\,$ This function works great for reading and writing data
- from flat files (delimited text files).
 vroom automatically uses the full abilities of your CPU, namely executes in multi-threaded (using the several processer cores that modern CPUs have).
- Let's install the vroom package and time writing the same data frame, let's call the file temp3.
 - o Note that vroom is using tab delimiter, so you need to specify delim=",".

- o If some R package is already installed, you could actually access a package function with :: without loading the full package with library() or require().
- system.time(vroom::vroom_write(a,"temp3.csv",d
 elim=",")

- Importing *other data types* directly (MS <u>Excel</u>, <u>SAS</u>, <u>SPSS</u>, <u>STATA</u>, <u>Minitab</u>, etc.) or importing data directly from *relational databases* is also available.
- Besides *vroom*, there are also dedicated packages to handle importing of flat file data.

 \circ For instance: *readr* and *data.table*.

• If you want to study any of the above in more details, please refer to:

https://www.datacamp.com/courses/importing-datain-r-part-1

https://www.datacamp.com/courses/importing-datain-r-part-2

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- A seemingly great free DataCamp tutorial gives details for reading different types of data is also available.
 - "... comprehensive, yet easy tutorial to quickly import data into R, going from simple text files to the more advanced SPSS and SAS files."

https://www.datacamp.com/community/tutorials
/r-data-import-tutorial

- Yet again, there is **rarely a real need to go fancy**.
 - \circ Just use a flat file format like . $\tt csv$ and move ahead with your project.
 - In my experience in almost all kinds of settings it is <u>almost always</u> better to use the KISS principle (credit: U.S. Navy in the mid-last century).

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- What about importing a real data set from the web?
- The **Stock Exchange of Hong Kong** is one of the fastest growing stock exchanges in Asia. It has 2,500+ listed companies with a combined market capitalization of more than 5 trillion USD.
 - \circ Let us consider an actual data set that is produced daily by this stock exchange.

 \circ Most files are flat tables.

 However, you often need to study the file before you are able to actually parse it in R (or any other language) ...

- The website is <u>https://www.hkex.com.hk/</u> o Or just google HK exchange.
- To save time we will download

https://www.hkex.com.hk/eng/dwrc/search/dwFullList.csv

 \circ Let's try directly downloading this $\tt csv$ file, supposedly comma separated values file.

 \circ What now?

- You need to look at the file or read the documentation...
- Small files like this, you can just download and investigate.
- Now let's resolve the issue...

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19. Simulations and Central Limit Theorem

• One of the true powers of R is to run simulations, **numeric experiments with random outcomes**, by sampling from:

the built-in R distributions
other distributions in external libraries/packages
or from the data directly.

- Recall, random sample from some distribution is done with *r*... where ... is replaced with the distribution name
 - For instance, rnorm(30) will generate 30 numbers from the standard normal distribution.

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• If you want to sample from data, one of the main tools is the sample() function.

sample(x,k) generates a random permutation
of k unique elements from the vector x, no
repeated elements are allowed by default

• You might want to allow repeats.

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• Allowing repetitions is like randomly selecting an element and "returning it back" in the set before the next selection (*sampling with replacement*).

sample(x,6,replace=T) # this works just fine

- Let us simulate rolling a fair die 6,000,000 times.
 - \circ Let's first "simulate" manually with the participants in our class session.
 - Randomly pick one integer from 1 to 6 and write it down.
 - Bar-graph (with the histogram or the barplot R function) is the default choice for a graphical representation here for non-continuous data.

- Now use R to simulate the same number of die rolls; store the results in rolls.
- We expect ~1/6 of the outcomes to be 4s. Let's compute how many we've got.
- How about 6,000,000 rolls in rolls?
- Let's seed the random generator with set.seed (2024) and compute how many of the 6,000,000 rolls are 4s.
- \bullet rolls vector is only about ${\sim}0.02Gb$ in the RAM. $_{\odot}$ Piece of cake for machines nowadays.

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- Let's estimate the **probability of getting** <u>at least one 6</u> in five consecutive rolls of a fair die.
 - \circ The exact (theoretical) probability is easy to derive...
 - We will use a quick simulation to "check" our theoretical result (computing experimental probability).
 - We should be very close to the theoretical probability if we have enough simulated die rolls.

Probability of getting at least one 6 in five consecutive die rolls...

• We will use the matrix function to store the results (although a loop can be used too).

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• Each column in the matrix will be a "trial", i.e., 5 die rolls.

```
# Always start modest when building code
trls <- 10  # only 10 die rolls
M <- matrix(sample(1:6,5*trls, replace=T),
ncol=trls)
# let's build our code one step a time ...
# ...
# Next, let's try 1,000,000 rolls
trls <- 10^6  # number of trials</pre>
```

The theoretical probability was...

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• Sometimes theoretical results are not feasible...

- Obtaining a theoretical **confidence interval for the ratio of two proportions** is such an example.
 - \circ The distribution of a ratio of random variables does not generally follow a simple known distribution.
 - In practice, we often use simulations to estimate the confidence interval for the ratio of two proportions, as exact theoretical results are difficult to obtain.
 - These methods are based on resampling techniques and can provide good approximations when the simulation size is large enough.

• Let's consider an example like this:

Suppose that in a survey of 250 males and 250 females, you observe that 88% of the females have health insurance, but only 80% of the males have it.

Computing a confidence interval for the ratio of these two proportions could be challenging but is needed as the data is just a sample.

Let us simulate to estimate the confidence interval ...

 Let us create a code to practice and to also illustrate the Central Limit Theorem Start from real data and create a simulated population following the instructions closely. We will do more than simply resampling 	<pre># Data source: data1 <- c(775, 900, 775, 820, 810, 1150, 790, 610, 875, 775, 890, 600, 760, 625, 710, 960, 500, 690, 745, 550, 685, 625, 775, 580, 590, 495, 595, 550, 495, 650, 580, 700, 1000, 900, 605, 475, 875, 850, 650, 785, 600, 825, 490, 500, 795)</pre>
# Analyzing Rental Data	# Instructions:
# This assignment will have you analyze rental price data # and take samples to compare sample means.	# 1. Generate a histogram of the 45 actual rental prices
# The data gives 45 one-bedroom apartment rental prices # listed on 1/18/2017 at rent.com and craigslist.	# 2. Create a total "population" of size 1000 Generate additional rental prices by:

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- Setting a seed for reproducibility: set.seed(12347)

- # Sampling from data1 with replacement
- # Adding random noise by sampling from normal # distribution centered around each sampled
- # value with SD=20. Each random number will
- # have a different mean, randomly picked from
 # the data.
- # Rounding the simulated values to nearest 10
- # Combing simulated and original rents in a
- # single vector aptRentPop

3. Calculate summary statistics of the population

4. Take 2 random samples of size 30 from the "population" in aptRentPop.

5. Calculate and compare the means of the two samples.

6. Take 1000 samples of size 30 from the "population" and store the 1000 sample means.

7. Calculate mean and standard deviation of the 1000 sample means.

8. Generate a histogram of the 1000 sample means.

9. Repeat the 1000 sample means to be for samples of size 9 this time.

Note that the original distribution was not close # to the normal distribution(not bell shaped). # The distribution of the sample means however is # approximately normal

This is the Central Limit Theorem in action

Central Limit Theorem (CLT)

The distribution of the sample mean is a probability distribution of all possible sample means for a given sample size.

- **CLT**: for a "large enough" sample size the **distribution of** the sample mean is approximately normal regardless of the distribution of the population we sample from.
- In our example, regardless of the shape of our "population" (aptRentPop), the distribution of the sample means tends toward normal distribution as the sample size increases.

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- **CLT**: For **any population**, the distribution of sample means is **approximately normal** for large enough sample size.
 - For large enough sample size the sample means have a distribution that can be approximated by a normal distribution with the population mean μ and population standard deviation $\frac{\sigma}{\sqrt{n}}$.
 - This assumes simple random sampling.
 - The concept of 'large enough' (often cited as a sample size greater than 30) varies depending on the distribution of the population.
 - o Different sample sizes are needed to achieve approximate normality distribution of the sample mean depending on the original population distribution.



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Skewed