## **Scientific Computation**

**Spring 2019** 

**Lecture 4** 

# **Today**

Constant-time search

- Lecture 2: Binary search applied to sorted array requires log<sub>2</sub>N operations
- Can we do better?

- Lecture 2: Binary search applied to sorted array requires log<sub>2</sub>N operations
- Can we do better?
- Proposal: Set the value of an integer as index in an array

6 8 23 32

- Then for the example above we would have an array (or list), X, where X[6] = X[8] = X[23] = X[32] = 1
- Speed for search would be constant time (O(1))
- But many weaknesses:
  - What about more general data (not just non-negative integers)
  - Inserting and deleting entries would require O(N) time

- Lecture 1: Binary search applied to sorted array requires log<sub>2</sub>N operations
- Can we do better?

- Real-world problem:
  - A startup is keeping track of unique visitors to its website each month
  - Each visitor must be assessed as "new" or "repeat"
  - Visitors who have not visited in 30+ days must be removed
  - Everything must be fast and accurate!

- Lecture 1: Binary search applied to sorted array requires log<sub>2</sub>N operations
- Can we do better?

- Visitors are identified by IP addresses
  - IP addresses are 4 8-bit numbers which (in base-10) take the form: 251.31.241.80 and identify a user's location on the internet
  - There are 256<sup>4</sup> possible addresses so it's not feasible to maintain an array for all possible addresses
  - Could maintain a sorted list but inserting new addresses would require O(N) operations (in Python)

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  - Could maintain a sorted list but inserting new addresses would require O(N) operations (in Python)
- Alternate approach: IP addresses are assigned indices based on their values
  - A hash function takes an address as input and provides and index as output
  - The design of hash functions is an important and active subject
  - What are the "desirable" properties of a hash function?

- Consider a hash function i = H(a)
  - i is a non-negative integer
  - a is a numeric representation of an IP address
  - Let's say we expect about 1e3 unique visitors each month
- Then, H should distribute these visitors to about 1e3 indices
  - Desirable property 1: It should always assign a particular IP to the same index on repeat visits

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- Then, H should distribute these visitors to about 1e3 indices
  - Desirable property 1: It should always assign a particular IP to the same index on repeat visits
- Since there are 256<sup>4</sup> possible IPs, it is impossible to design H so that unique visitors are always assigned unique indices
  - When two or more different IPs are assigned the same index, we have a hash collision
  - Say 0<=i<=N. Desirable property 2: Probability of 2 distinct visitors being assigned same index, i should be 1/N

#### An example:

- The IP address is four integers, a<sub>1</sub>,a<sub>2</sub>,a<sub>3</sub>,a<sub>4</sub>
- Choose four arbitrary integer weights, w<sub>1</sub>,w<sub>2</sub>,w<sub>3</sub>,w<sub>4</sub>
- Is:  $i = \sum w_i a_i$  a suitable hash function?

#### An example:

- The IP address is four integers, a<sub>1</sub>,a<sub>2</sub>,a<sub>3</sub>,a<sub>4</sub>
- Randomly choose four arbitrary integer weights, w<sub>1</sub>,w<sub>2</sub>,w<sub>3</sub>,w<sub>4</sub>
- Is:  $i = \sum w_i a_i$  a suitable hash function?
- Not quite:
  - This can generate 2564 indices, we only want about 1e3
- What about  $i = (\Sigma w_i a_i) \mod 1000$ ?
  - We'll now have the right number of indices
  - But if there are patterns in the IPs, could have large number of visitors assigned the same index
- Better to use a prime number,  $i = \sum w_i a_i \mod 997$

Better to use a prime number,  $i = h(a_j) = \sum w_j a_j \mod 997$ 

- How well does this function work?
- Given two different addresses a and b, what is the probability that h(a) = h(b)?

Let's assume that  $a_4 \neq b_4$  and that the weights are 4 random integers mod 997

• There will be a hash collision if:  $\sum_{j=1}^4 w_j a_j \equiv \sum_{j=1}^4 w_j b_j \pmod{997}$ 

where  $x \equiv y \pmod{n}$  if n divides x - y

- Rearranging:  $\sum_{j=1}^{3} w_j (a_j b_j) \equiv w_4 (b_4 a_4) \pmod{997}$
- The LHS is just some, integer, c, and we need to "solve" the following congruence for  $w_4$ :  $w_4(b_4-a_4) \equiv c \pmod{997}$

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#### To solve this for $W_4$ , need a result from modular arithmetic:

Let p be a prime. If k is not a multiple of p, then there exists an integer  $k^{-1} \in \{1, 2, ..., p-1\}$  such that:  $k \cdot k^{-1} \equiv 1 \pmod{p}$ 

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In our example, p=997, k = (b<sub>4</sub>-a<sub>4</sub>) so, 
$$w_4 \equiv c(b_4 - a_4)^{-1} \pmod{997}$$

We *choose:*  $w_4 = c(b_4-a_4)^{-1} \mod 997$  and more generally, require:  $w_i \in \{0, 1, ..., p-1\}$ 

This allow us to state that there is only one  $w_4$  in the range [1 p-1] for which a and b will be assigned the same index. So, probability(h(a)=h(b)) = 1/p.

- We now have a hash function that will assign indices to IP addresses
- What is the general workflow?
  - 0. Initialize a dictionary or list (a *hash table*) where you will store addresses
  - 1. Given an IP, compute an index
  - 2. And store the IP in the corresponding location in the hash table
  - 3. Append the IP if there is already an address in the computed list

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- Let's assume that we have M IP addresses and N indices with M>N
- On average, each index will be assigned M/N IPs
- Should choose M to be a prime number close to N (in practice, maybe close to 2N)

We started looking for something faster than binary search

What is the cost of using a hash table?

- Search (lookup): Given an IP, find it in the table
  - Evaluate the hash function and obtain an index
  - Check the number of items stored at index, i.
  - If i > 1, iterate through items
  - Overall, O(max(1,M/N))
  - For well-designed hash table and function, will be close to O(1)!
- Insert: Add an IP to table
  - Search
  - Then append at computed index if IP is not already present
  - Overall, close to O(1) for a careful implementation
- Delete: Remove an IP
  - Same as Insert, just replace "append" with "delete"

#### **Summary:**

- Binary search: O(log<sub>2</sub>N) but requires the maintenance of a sorted list/array
- Hash table: O(1) for search as well as maintenance
  - Provided that the hash function and table are both welldesigned!

#### **Summary:**

- Binary search: O(log<sub>2</sub>N) but requires the maintenance of a sorted list/array
- Hash table: O(1) for search as well as maintenance
  - Provided that the hash function and table are both welldesigned!
- What does this look like in Python?
  - Let's move away from the IP problem to a more general task
  - Consider input that may be real numbers or even non-numeric

## **Python hash function**

Python provides a hash function that returns an integer for (almost) any input

- For an integer, i, hash(i) = i
- For two inputs if a==b then hash(a)=hash(b)
  - hash(3.0) = hash(3) = 3
- For non-integers, the function is less predictable:

```
In [1]: hash('m3sc')
Out[1]: 5237192937700339269
In [2]: hash('m3c')
Out[2]: 1091467703030128764
```

## Python hash table

#### Can use list of lists to build hash table

- Compute hash for each item of interest
- Store item using hash value as index for list
- For hash collisions, append item at index
- Lookup:
  - Compute hash value
  - Search through sub-list at index for item
- Not too difficult try it!

- But we don't have to build our own hash table!
- Python dictionaries are hash tables (technically, they are "associative arrays")
- Dictionaries are containers where each element is a key-value pair
  - A key can be considered a label or id

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  - A key can be considered a label or id

 Python applies a hash function to keys to know where to store them and where to look for them

• Important dictionary operations Constant time O(1):

```
In [36]: d = dict()  #initialize a new dictionary
In [37]: d[key] = value #associate key with value and store in d
In [38]: d[key]  #value associated with key in d, raises KeyError if key has not been added to d
Out[38]: [14, 1, 2019, 20]
In [39]: x=1
In [40]: d.get(key,x)  #value associated with key if key is present,
```

otherwise x

• Important dictionary operations Constant time O(1):

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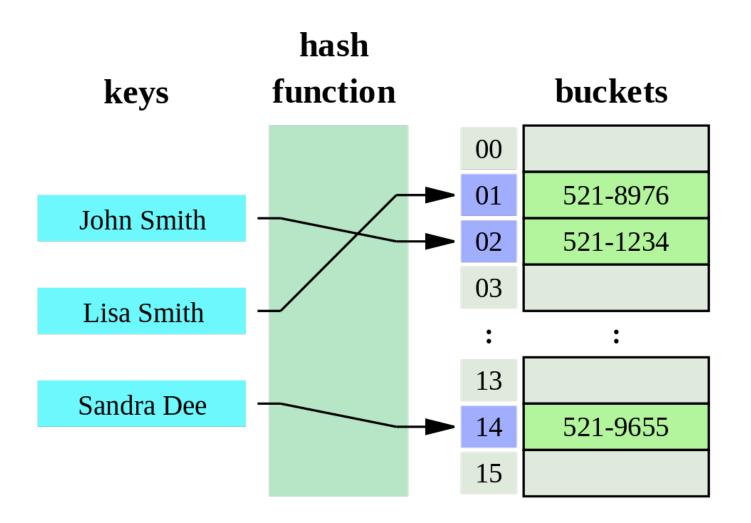
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In [40]: d.get(key,x) #value associated with key if key is present,
otherwise x
In [42]: key in d #is key in d?
Out[42]: True
In [43]: len(d) #number of key-value pairs in d
Out [43]: 1
In [44]: del d[key] #remove key (and its associated value) from d
```

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In [44]: del d[key] #remove key (and its associated value) from d
linear time O(N): In [45]: for key in d: # iterate over keys in d
```

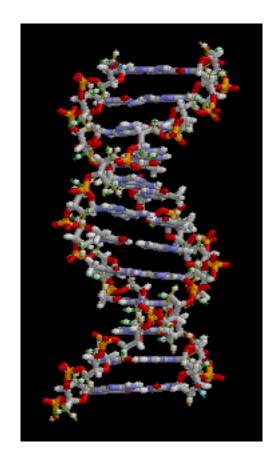
This is exactly what we need to build and maintain a hash table.

## Python hash table



### Genetic code

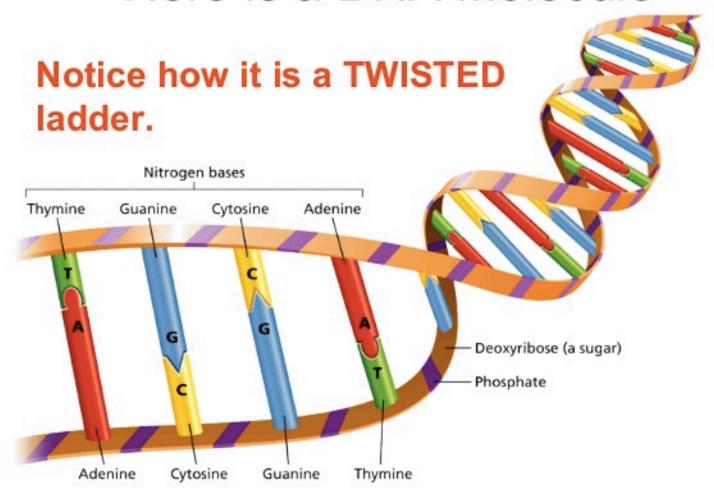
- DNA (and RNA) consists of two "strands" arranged in a double helix and connected with covalent bonds
- Each strand contains a sequence of nucleotides
- DNA is constructed from 4 nucleotides:
  - Adenine
  - Cytosine
  - Guanine
  - Thymine (RNA has Uracil in place of Thymine)



### Genetic code

- DNA is constructed from 4 nucleotides (or bases):
  - Adenine
  - Cytosine
  - Guanine
  - Thymine (RNA has Uracil in place of Thymine)
- Adenine bonds with Thymine and Guanine bonds with Cytosine
- So if one strand contains the sequence GCTTCA the other strand will contain CGAAGT in the corresponding location
- During cell division, each daughter cell gets one strand
  - And then the needed 2<sup>nd</sup> strand can be constructed so A's pair with T's and C's pair with G's

## Here is a DNA Molecule



### Genetic code

- DNA is constructed from 4 nucleotides (or bases):
  - Adenine
  - Cytosine
  - Guanine
  - Thymine (RNA has Uracil in place of Thymine)
- Codons consist of three DNA bases and contain code for synthesizing amino acids
  - Proteins are built from amino acids
  - 64 possible codons, but there are only 20 essential amino acids specified by DNA
- Gene sequencing involves:
  - Extracting the sequence of bases from DNA samples
  - Investigating the proteins or functions associated with codons and their sequences

## Pattern finding

- A fundamental computational problem:
  - Given a DNA sequence, search for a pattern
  - Both the sequence and the pattern can be extremely long
  - The number of patterns can also be large
- Fruit fly genome: 139.5 million base pairs



Humans: 3 billion or 6 billion pairs (depending on the cell type)



How can we efficiently search for patterns?

### Pattern search

Problem setup:

Specify a N-character sequence, S, and a M-character pattern, P Find all locations in S where P occurs

Example:

S = ATGTTGTACCGTATCGG P = GTA

N=16, M=3