Randomised Algorithms

How many statisticians does it take to screw in a light bulb?

One. But then you're only 95% confident it's been changed.

Most algorithms are deterministic. If given the same input, they will return the same output. Every time.

A randomised algorithm will return different outputs for the same input.

That might seem weird: if there is one optimum value for a problem, then shouldn't I always return that?

Well, in most practical optimisation problems we actually want to find a *good* solution. The *best* would be icing, but sometimes that would require almost infinitely more computation to find (that isn't hyperbola).

There are very many examples of randomised algorithms:

- · You will see some applied to optimisation in this course.
- Statistical sampling is an almost too obvious case.
- The general class of MCMC for all sorts of stuff.
- Some neural network algorithms start from a random set of weights.

I want to cover some examples you might not normally see in a maths course.

Key example: random hashes

Hash Functions

Imagine my data objects $x \in \Omega$ (the set Ω is sometimes called the *universe*, and the objects the *keys*). The objects can be completely unstructured (or not).

A hash function $h:\Omega\to [m]=\{0,\ldots,m-1\}$. That is, it is a function that maps the objects x to the set $\{0,\ldots,m-1\}$ denoted by [m]. We often call the elements of the set bins Usually the size of the potential input universe $|\Omega|\gg m$.

Most hash functions are many-to-one. Many hash functions are *surjective* on non-trivial data sets, that is, for every $i\in[m]$ there is at least one potential input element $x\in\Omega$ such that h(x)=i. But hash functions are not, in general, invertable. That is, you can't use h(x) to work out x.

A *collision* occurs if h(x) = h(y) for $x \neq y$. That is, two different inputs are mapped to the same hash value.

Hash functions often occur in families H, sometimes indexed by a parameter, e.g., we might write $H = \{h_i\}_{i=1}^n$. The index set is often assumed to be very large for cryptography applications, but we don't need that here.

Desirable Properties of Hash Functions

Hash functions have a number of desirable properties depending on application.

- In most cases it is important that they be either fast, or very, very fast.
- We would like the hash function to spread out the data roughly evenly over [m] to avoid collisions as much as possible.

Random Hash Functions

Input data is not really random, so often we use families of random functions, particularly in order to define properties in a formal sense.

In this context, uniformity can be defined by

$$P_{h\in H}ig(h(x)=i)ig)=rac{1}{m},$$

for $x\in\Omega$ and $i\in[m]$. Remember that x is not random here, the function h is. Uniformity is appealing, but can be achieved by stupid hash functions such as the constant $h_i(x)=i$.

A family of hash functions is called *universal* if it satisfies the property that for all $x,y\in\Omega$ with x
eq y

$$P_{h \in H} \leq h(y) \log \frac{1}{m}.$$

That says that two different inputs collide with a probability at most 1/m when we draw a random hash function from H.

Universality might be hard to achieve, particularly with a fast hash, so we often allow near-universality, by putting a constant factor (say of 2) into the above.

Examples

- 1. **Linear algebra**: if the inputs are vectors $\mathbf{x} \in \{0,1\}^n$ (0-1 vectors) then we can perform hashing through matrix multiplication by an $m \times n$ 0-1 matrix. We can construct a family of hashes by taking random 0-1 matrices. This example is illustrative but rarely used, because it only works for very restrictive data. More generally we would like hashes that map unstructured data.
- 2. **Modular arithmetic:** Take $h_i(x) = x + i \mod m$. Or do some multiplication and addition first to get

$$h_{a,b}(x) = ((ax+b) \mod p) \mod m,$$

for some given (fixed) prime number p. Chose the right p and set of a and b you get a universal hash family. But this only works for input integers.

2. **Recursive hashes:** You can take any input of a fixed length and convert it first to an integer as a block, and then use modular arithmetic to convert it. But what do you do with arbitrary length data like natural language text? You could pad the text out to a fixed length, but that might have to be really large. You can instead treat the string like a polynomial and use a recursive hash, which is really Horner's method for computing polynomials, but in general this looks like:

$$h_a(\mathrm{'xyf'}) = a \oplus h_x(\mathrm{'yf'}).$$

This is commonly called a rolling hash.

Recursion is a general strategy for dealing with fast data – the cost for each element is a small constant.

In many ways such hash function are linked to pseudo-random number generation as well.

The above aren't stupidly slow, but there are faster hashes. If we can use only base-2 arithmetic (multiplication and division by 2) we can make these much faster, for instance.

Applications

A core use for hashes in data science is to enable fast indexing of a data set. That is, a fast way to look up, or add data to a larger repository of potentially unstructured data. A "hash table" is a standard data structure used in most modern programming languages to create associative arrays (also called hashes, and dictionaries). Naively you just store x in a fixed array at location h(x), but it is a little more complicated than that.

But they have many other uses. A key one is in cryptography, but I won't talk about that here. What I will talk about is some cute ways to summarise data very, very quickly to deal with fast data.

Calculating Similarity of Two Giant Sets

Imagine two very, very large sets A and B. A common measure of similarity (roughly the opposite of distance) between sets is the $Jaccard\ Index$

$$J(A,B) = rac{A \cap B}{A \cup B}$$

Imagine, for instance, that A and B are the sets of words that appear in two documents. If the two are identical, then J(A,B)=1. If they are completely different, then J(A,B)=0. Mostly likely they are somewhere in between.

A naive algorithm would check, for every element $a\in A$ if that element also appeared in B. That would take $O(|A|\times |B|)$ to do the dumb way. We could use hash tables to speed it up. But there is a fast approximate approach called MinHash.

The **MinHash** algorithm provides a fast means calculate an approximation to J(A,B). Its usually described in terms of random permutations in order to prove its properties, but the standard implementation is as follows

Or at least my best guess is:

Start with sets S_j to compare, with union Ω and one main hash $g:\Omega\to\{1,\ldots,N\}$. Take k random hash functions $\{h_1,\ldots,h_k\}$ and initialise k values $\{c_1,\ldots,c_k\}$ to ∞ . We create a signature of a set as follows:

- 1. We start by mapping the elements of the sets we want to compare to numbers $\{1,\dots,N\}$ using a hash f.
- 2. Use a hash table g(S) to index all of the elements of S
- 3. For all $x \in \Omega$ (or for i=1 to N)

If
$$g(x) \in g(S)$$
 then take $i = g(x)$

For
$$i = 1$$
 to k

If
$$h_i(i) < c_i$$
 then $c_i \leftarrow h_i(i)$

The signature of S_j is the vector $\mathbf{m}(S_j) = \mathbf{c}$.

The Jaccard index is computed by taking

$$J(A,B) = \frac{1}{k} \sum_{i=1}^{k} I(m_j(A) = m_j(B)),$$

where $I(\cdot)$ is an indicator function. Intuitively this is because the probability that $m_j(A)=m_j(B)$ for any given random hash h_j is proportional to the Jaccard index of the two sets.

The accuracy of the approach depends on k, but the computation is proportional to k, so there is a tradeoff.

There are very many algorithms in this space, showing how hashes can be used (particularly groups of random hashes) to construct various approximate summaries of a data set.

- **Bloom filters** let you quickly test if an element $x \in S$. False positives are possible, but false negatives are not.
- The **Count Min** sketch lets you count (approximately) the number of times an element appears in
- We can use hashes (in *feature hashing* or the *hashing trick*) to vectorise a set of features, i.e., put them into tensor form.

They can often be applied to rapidly growing data sets.

To do (1 of the the following)

In Python (using PyTorch or not)

- 1. Write a rolling hash function for images. Use it to hash the MNIST images into 6 bins. Check the uniformity of your hash on this data.
- 2. Implement MinHash. Generate two largish random sets of integers from 1-N (avoid the first hashing step), and then compute their Jaccard index both directly and using MinHash.

See jaccard.ipynb

Terminology

Lookup the following:

- FPRAS
- · Monte-Carlo vs Los-Vegas Algorithms
- PAC learning
- VC-dimension

There are many similarity indexes that are similar to Jaccard, e.g., the Dice coefficient.

Links

- https://brilliant.org/wiki/randomized-algorithms-overview/
- http://theory.stanford.edu/people/pragh/amstalk.pdf
- https://datascience.stackexchange.com/questions/32557/what-is-the-exact-definition-of-vc-dimension
- https://www.cs.utah.edu/~zhe/pdf/lec-13-pac-definition-upload.pdf
- https://www.cs.princeton.edu/courses/archive/spr08/cos511/scribe_notes/0211.pdf
- https://jeremykun.com/2014/01/02/probably-approximately-correct-a-formal-theory-of-learning/
- https://courses.engr.illinois.edu/cs473/sp2017/notes/05-hashing.pdf
- https://ageek.dev/polynomial-rolling-hash
- https://www.cs.cmu.edu/afs/cs/project/pscico-guyb/realworld/www/slidesS14/minhash.pdf
- https://florian.github.io/count-min-sketch/
- http://dimacs.rutgers.edu/~graham/pubs/papers/cmencyc.pdf
- https://en.wikipedia.org/wiki/Feature_hashing