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Fast Fourier Transform and 2D Convolutions

Doing convolutions fast

Stephen Huan¹

¹Thomas Jefferson High School for Science and Technology

TJ Vision & Graphics Club, October 21, 2020

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The Fast Fourier Transform (FFT) is a common technique for signal processing and has many engineering applications. It also has a fairly deep mathematical basis, but we will ignore both those angles in favor of accessibility. Instead, we will approach the FFT from the most intuitive angle, **polynomial multiplication**.

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The Fast Fourier Transform (FFT) is a common technique for signal processing and has many engineering applications. It also has a fairly deep mathematical basis, but we will ignore both those angles in favor of accessibility. Instead, we will approach the FFT from the most intuitive angle, **polynomial multiplication**.

First, we represent polynomials by a list of coefficients, where the number at index 0 represents the coefficient of x^0 , the number at index 1 represents the coefficient of x^1 , and so on.

Polynomial representation

The polynomial $3 + 2x + 4x^2$ becomes $[3, 2, 4]$.

Definition of the Convolution

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The multiplication of two polynomials f and g is then simply each term of f multiplied with each term of g and then added up. We can also assume that f and g are the same length N , where the polynomial of lesser degree is padded with zeros. If we say the product is p , we can give an formula for an index in p in the following way:

Definition of the convolution

$$p[n] = (f * g)[n] = \sum_{i=0}^n f[i]g[n-i]$$

Intuition Behind a Convolution

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$p[n]$ is the coefficient of x^n in the product, and it is formed by adding up all the possible ways to get to x^n , i.e. $f[0]x^0$ times $g[n]x^n$, $f[1]x^1$ times $g[n-1]x^{n-1}$, etc. Intuitively, this “flips” g , and then the resulting product is computed by “sliding” g over f and then computing the dot product between the two lists, or a weighted average.

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Suppose we have the polynomial $3 + 2x + 4x^2 = [3, 2, 4]$ and the polynomial $1 + 3x + 2x^2 = [1, 3, 2]$. To compute their product, we first flip the second list to get $[2, 3, 1]$.

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Suppose we have the polynomial $3 + 2x + 4x^2 = [3, 2, 4]$ and the polynomial $1 + 3x + 2x^2 = [1, 3, 2]$. To compute their product, we first flip the second list to get $[2, 3, 1]$. We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

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We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

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We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

Then, $[3, 1]$ overlaps with $[3, 2]$ so we get $3 \cdot 3 + 1 \cdot 2 = 11$.

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We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

Then, $[3, 1]$ overlaps with $[3, 2]$ so we get $3 \cdot 3 + 1 \cdot 2 = 11$.

$[2, 3, 1]$ overlaps with $[3, 2, 4] = 16$

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We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

Then, $[3, 1]$ overlaps with $[3, 2]$ so we get $3 \cdot 3 + 1 \cdot 2 = 11$.

$[2, 3, 1]$ overlaps with $[3, 2, 4] = 16$

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We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

Then, $[3, 1]$ overlaps with $[3, 2]$ so we get $3 \cdot 3 + 1 \cdot 2 = 11$.

$[2, 3, 1]$ overlaps with $[3, 2, 4] = 16$

$[2, 3]$ overlaps $[2, 4] = 16$,

and finally $[2]$ overlaps with $[4]$ to give 8.

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Suppose we have the polynomial $3 + 2x + 4x^2 = [3, 2, 4]$ and the polynomial $1 + 3x + 2x^2 = [1, 3, 2]$. To compute their product, we first flip the second list to get $[2, 3, 1]$.

We then slide $[2, 3, 1]$ over $[3, 2, 4]$, imagining there are zeros such that the parts of $[2, 3, 1]$ that don't overlap with $[3, 2, 4]$ aren't counted.

For the first value, 1 overlaps with 3 so we get 3.

Then, $[3, 1]$ overlaps with $[3, 2]$ so we get $3 \cdot 3 + 1 \cdot 2 = 11$.

$[2, 3, 1]$ overlaps with $[3, 2, 4] = 16$

$[2, 3]$ overlaps $[2, 4] = 16$,

and finally $[2]$ overlaps with $[4]$ to give 8.

Our final answer is then $[3, 11, 16, 16, 8] =$

$3 + 11x + 16x^2 + 16x^3 + 8x^4 = (3 + 2x + 4x^2)(1 + 3x + 2x^2)$.

Commutativity of the Convolution

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What if we computed $g * f$? It should be the same since polynomial multiplication should be commutative.

Theorem

$f * g = g * f$, i.e. *polynomial multiplication is commutative.*

Proof of the Commutativity of the Convolution

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Proof.

We have $(f * g)(n) = \sum_{i=0}^n f[i]g[n-i]$ by definition. Perform the variable substitution $k = n - i$, so $i = n - k$. Summing from $\sum_{i=0}^n$ will sum from $k = n$ to $k = 0$ in descending order, so $\sum_{i=0}^n = \sum_{k=0}^n$ (from the commutativity of addition).

$$(f * g)[n] = \sum_{i=0}^n f[i]g[n-i] \quad \text{Definition}$$

$$= \sum_{k=0}^n f[n-k]g[k] \quad \text{Substitution}$$

$$= (g * f)$$



The Fast Fourier Transform

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This operation is known as a **convolution**, which is equivalent to polynomial multiplication in the discrete case and is denoted $f * g$. Its relevance to image processing will be expounded on later (for now, this puts the “convolutional” in “Convolutional Neural Networks”). Today’s lecture is about the **Fast Fourier Transform**, an efficient algorithm to perform convolutions.

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A naive approach to the convolution of two lists of length N, M will have runtime $O(NM)$ using the standard polynomial multiplication algorithm (each term of the first list multiplied with each term of the second list).

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A naive approach to the convolution of two lists of length N, M will have runtime $O(NM)$ using the standard polynomial multiplication algorithm (each term of the first list multiplied with each term of the second list).

What will be the length of the resulting list?

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A naive approach to the convolution of two lists of length N, M will have runtime $O(NM)$ using the standard polynomial multiplication algorithm (each term of the first list multiplied with each term of the second list).

What will be the length of the resulting list?

The first list is a polynomial of degree $N - 1$, the second of degree $M - 1$, so the resulting polynomial has degree $(N - 1) + (M - 1) = N + M - 2$.

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A polynomial of degree D has $D + 1$ coefficients, so the length of the product is $N + M - 1$. Thus,

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A polynomial of degree D has $D + 1$ coefficients, so the length of the product is $N + M - 1$. Thus,

Lower bound for the runtime of a convolution

The length of the convolution is $N + M - 1$, which is linear, so a better runtime than quadratic could exist.

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Point-Value Representation

An alternative way to express a polynomial

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The key observation is that we can represent polynomials in a different form than a coefficient list. In particular, we can use a **point-value** representation, or a list of (x, y) pairs that give an input and the corresponding output of a polynomial.

Point-value representation

A point-value representation for a polynomial p is a list of distinct x values and their corresponding y values, e.g.

$$\{(x_0, p(x_0)), (x_1, p(x_1)), \dots, (x_n, p(x_n))\}$$

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Evaluation

We call the process of going from a coefficient representation to a point-value representation ***evaluation***, since we evaluate the polynomial at multiple points to get the point-value representation.

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Evaluation

We call the process of going from a coefficient representation to a point-value representation ***evaluation***, since we evaluate the polynomial at multiple points to get the point-value representation.

Interpolation

Likewise, we call the process of going from a point-value representation to a coefficient representation ***interpolation***, since we are finding a polynomial which “fits” the data.

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Evaluation

We call the process of going from a coefficient representation to a point-value representation ***evaluation***, since we evaluate the polynomial at multiple points to get the point-value representation.

Interpolation

Likewise, we call the process of going from a point-value representation to a coefficient representation ***interpolation***, since we are finding a polynomial which “fits” the data.

Suppose we have a polynomial of degree n . We then need a certain number of points for evaluation and interpolation to be well-defined.

Well-defined Representations

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Evaluation is always well-defined, because we can always evaluate a polynomial of any degree or coefficient representation. However, if we don't have enough points, interpolation is not necessarily possible.

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Evaluation is always well-defined, because we can always evaluate a polynomial of any degree or coefficient representation. However, if we don't have enough points, interpolation is not necessarily possible.

Ill-defined representation

Consider the point-value representation $[(0, 0), (1, 1)]$ and a degree of 2. This could be the polynomial x^2 or $2x^2 - x$.

Well-defined Representations

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Evaluation is always well-defined, because we can always evaluate a polynomial of any degree or coefficient representation. However, if we don't have enough points, interpolation is not necessarily possible.

Ill-defined representation

Consider the point-value representation $[(0, 0), (1, 1)]$ and a degree of 2. This could be the polynomial x^2 or $2x^2 - x$.

So for a polynomial of degree n , we need at least $n + 1$ distinct points (since each point gives another linear equation constraining the $n + 1$ coefficients of the polynomial). We can in fact prove that if we have $n + 1$ points, that uniquely determines a polynomial of degree n .

Proof of the Point-Value Representation

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Theorem

A point-value representation with n distinct points uniquely determines a polynomial of degree $n - 1$.

Proof.

We have a polynomial of the form

$p(x) = a_0 + a_1x + a_2x^2 + \cdots + a_{n-1}x^{n-1}$ and n points of the form (x_i, y_i) such that $p(x_i) = y_i$. Those constraints determine the following matrix equation:

$$\begin{bmatrix} 1 & x_0 & x_0^2 & \cdots & x_0^{n-1} \\ 1 & x_1 & x_1^2 & \cdots & x_1^{n-1} \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ 1 & x_{n-1} & x_{n-1}^2 & \cdots & x_{n-1}^{n-1} \end{bmatrix} \begin{bmatrix} a_0 \\ a_1 \\ \vdots \\ a_{n-1} \end{bmatrix} = \begin{bmatrix} y_0 \\ y_1 \\ \vdots \\ y_{n-1} \end{bmatrix}$$

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Proof.

The leftmost matrix is known as the *Vandermonde matrix*, denoted $V(x_0, x_1, \dots, x_{n-1})$ which has the determinant (left as an exercise for the reader)

$$\prod_{0 \leq j < i \leq n-1} (x_i - x_j)$$

A matrix is invertible if and only if its determinant is nonzero, so this matrix is invertible if each x_i is distinct. Thus, we can solve for the coefficients by multiplying by the inverse, so $\vec{a} = V^{-1}\vec{y}$, and this solution is unique since an invertible matrix is a **bijective** transformation between a vector space and itself. \square

Lagrange's Formula

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This proof directly gave an easy construction of the interpolating polynomial, by $V^{-1}\vec{y}$. Matrix inverses can be computed in $O(n^3)$ as an easy upper bound, but that can be improved with **Lagrange's interpolating formula** to yield a $O(n^2)$ time algorithm. I will not elaborate on Lagrange's formula in this lecture but a good Wikipedia page is available [here](#).

Evaluation

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If we have a list of N coefficients, then the polynomial is of degree $N - 1$ and thus we need N distinct points. We first figure out how to evaluate polynomial at a single point, and will repeat the process for all the points.

Evaluating a Polynomial

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Suppose we have a polynomial of the form $p(x) = a_0 + a_1x + a_2x^2 + \dots + a_{n-1}x^{n-1}$. If we evaluate at a particular x_0 , we compute each $a_ix_0^i$ term which would take $O(N^2)$ time with repeated multiplication and $O(N \log N)$ time with fast exponentiation.

Horner's Rule

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But we can do better with **Horner's rule**. We notice that the degree in coefficient form is monotonically increasing, so we can successively factor out a multiplication by x .

Horner's rule

$$p(x) = a_0 + x(a_1 + x(a_2 + \cdots + x(a_{n-2} + xa_{n-1})))$$

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Horner's rule

$$p(x) = a_0 + x(a_1 + x(a_2 + \cdots + x(a_{n-2} + xa_{n-1})))$$

We do N multiplications and N additions, so the algorithm runs in $O(N)$. Evaluating a polynomial at N points then takes $O(N \cdot N) = O(N^2)$ time.

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So we can do both evaluation and interpolation in $O(n^2)$ and both are well-defined if we have enough points. Why did we figure this out? We can multiply two polynomials efficiently if we have the point-value representations of each!

Multiplying in Linear Time

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Suppose we have polynomials f, g in coefficient form. We also assume that the polynomials are evaluated at the *same* points, so we have

$$[(x_0, f(x_0)), (x_1, f(x_1)), \dots]$$

and

$$[(x_0, g(x_0)), (x_1, g(x_1)), \dots]$$

$f * g$ is then simply

$$[(x_0, f(x_0)g(x_0)), (x_1, f(x_1)g(x_1)), \dots]$$

or the element-wise multiplication of the two lists.

This can be easily computed in $O(n)$!

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So our algorithm for polynomial multiplication is as follows:

- 1 Evaluate a coefficient representation into a point-value representation.

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So our algorithm for polynomial multiplication is as follows:

- 1 Evaluate a coefficient representation into a point-value representation.
- 2 Multiply the two point-value representations in linear time.

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So our algorithm for polynomial multiplication is as follows:

- 1 Evaluate a coefficient representation into a point-value representation.
- 2 Multiply the two point-value representations in linear time.
- 3 Interpolate the resulting point-value representation back to coefficients.

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The speed of this algorithm is contingent on our ability to quickly evaluate and interpolate a polynomial. Currently, with our $O(n^2)$ time evaluation and interpolation algorithms we match the $O(n^2)$ naive algorithm. However, under this framework we can improve the time if we pick our points cleverly rather than arbitrarily.

Complex Roots of Unity

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Our special points are going to be **complex roots of unity**, or roots of 1 that are allowed to have an imaginary component.

Complex roots of unity

The second root of 1 can be 1 or -1 (taking “second root” to mean anything which squared is 1). The fourth root of 1 can be 1, -1 , i , or $-i$. (since $i^4 = (i^2)^2 = (-1)^2 = 1$).

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To easily compute these roots, we can rewrite 1 using

Euler's formula

$e^{ix} = \cos x + i \sin x$ (a proof of this appears in the appendix).

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To easily compute these roots, we can rewrite 1 using

Euler's formula

$e^{ix} = \cos x + i \sin x$ (a proof of this appears in the appendix).

$$e^{2\pi i} = \cos 2\pi + i \sin 2\pi = 1$$

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To easily compute these roots, we can rewrite 1 using

Euler's formula

$e^{ix} = \cos x + i \sin x$ (a proof of this appears in the appendix).

$$e^{2\pi i} = \cos 2\pi + i \sin 2\pi = 1$$

So we can take a n th root by simply raising

$$\sqrt[n]{1} = (e^{2\pi i})^{\frac{1}{n}}$$

so a root is

$$e^{\frac{2\pi i}{n}}$$

Properties of Complex Roots

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However, note that we can rewrite 1 in many different ways since sine and cosine are periodic.

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However, note that we can rewrite 1 in many different ways since sine and cosine are periodic.

Since adding 2π doesn't change the value of sine and cosine, 1 is also equal to $e^{4\pi i}$, $e^{6\pi i}$, and so on.

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However, note that we can rewrite 1 in many different ways since sine and cosine are periodic.

Since adding 2π doesn't change the value of sine and cosine, 1 is also equal to $e^{4\pi i}$, $e^{6\pi i}$, and so on.

In general, $e^{2\pi ki}$ is equal to 1 for any integer k , so if we take the n th root, $e^{\frac{2\pi ki}{n}}$ is also going to be a valid root of unity.

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Since adding 2π doesn't change the value of sine and cosine, 1 is also equal to $e^{4\pi i}$, $e^{6\pi i}$, and so on.

In general, $e^{2\pi ki}$ is equal to 1 for any integer k , so if we take the n th root, $e^{\frac{2\pi ki}{n}}$ is also going to be a valid root of unity.

However, not every k gives a distinct root of unity.

$k = n + 1$ is equivalent to $k = 1$ since

$$\cos\left(\frac{2\pi(n+1)}{n}\right) = \cos\left(2\pi + \frac{2\pi}{n}\right) = \cos\left(\frac{2\pi}{n}\right)$$

This generalizes such that an power k equivalent to $j \bmod n$ will have the same root.

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We can easily keep track of the n distinct n th roots of unity by writing them as powers of the **principle root of unity**.

Principle Root of Unity

The *principle root of unity* is the root of unity when $k = 1$.

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We can easily keep track of the n distinct n th roots of unity by writing them as powers of the **principle root of unity**.

Principle Root of Unity

The *principle root of unity* is the root of unity when $k = 1$.

Notation

We will denote this principle root as ω_n , where $\omega_n = e^{\frac{2\pi i}{n}}$.

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We can easily keep track of the n distinct n th roots of unity by writing them as powers of the **principle root of unity**.

Principle Root of Unity

The *principle root of unity* is the root of unity when $k = 1$.

Notation

We will denote this principle root as ω_n , where $\omega_n = e^{\frac{2\pi i}{n}}$.

Since we picked $k = 1$, we can represent every n th root of unity as a power of this root of unity since

$$e^{\frac{2\pi ki}{n}} = (e^{\frac{2\pi i}{n}})^k = \omega^k$$

Properties of a Principle Root

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Note that every power of the principle root of unity is itself a root of unity, because

$$(\omega_n^k)^n = (\omega_n^n)^k = 1^k = 1$$

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Note that every power of the principle root of unity is itself a root of unity, because

$$(\omega_n^k)^n = (\omega_n^n)^k = 1^k = 1$$

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We now come to an observation that will be instrumental in developing the FFT - that the square of a n th principle root of unity is a $\frac{n}{2}$ th principle root of unity.

This follows nearly from definition:

$$\begin{aligned}\omega_n^2 &= (e^{\frac{2\pi i}{n}})^2 \\ &= e^{\frac{4\pi i}{n}} \\ &= e^{\frac{2\pi i}{\frac{n}{2}}} \\ &= \omega_{\frac{n}{2}}\end{aligned}$$

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We now show that evaluating a polynomial at n distinct n th roots of unity can be written as a recurrence relation. Our observation is a clever rewrite of a polynomial into two parts.

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Suppose we have the polynomial

$$p(x) = a_0 + a_1x + a_2x^2 + \cdots + a_nx^{n-1}$$

We divide the coefficient list of p into two parts, one with even powers and the other with odd powers, the left and right halves respectively.

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Suppose we have the polynomial

$$p(x) = a_0 + a_1x + a_2x^2 + \cdots + a_nx^{n-1}$$

We divide the coefficient list of p into two parts, one with even powers and the other with odd powers, the left and right halves respectively.

We assume that n is a power of 2 so that p can always be divided in such a manner.

If n isn't, we can always pad with 0's.

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Suppose we have the polynomial

$$p(x) = a_0 + a_1x + a_2x^2 + \cdots + a_nx^{n-1}$$

We divide the coefficient list of p into two parts, one with even powers and the other with odd powers, the left and right halves respectively.

We assume that n is a power of 2 so that p can always be divided in such a manner.

If n isn't, we can always pad with 0's.

$$p(x) = a_0 + a_1x + a_2x^2 + \cdots + a_nx^{n-1} \quad (1)$$

$$L(x) = a_0 + a_2x + a_4x^2 + \cdots \quad (2)$$

$$R(x) = a_1 + a_3x + a_5x^2 + \cdots \quad (3)$$

It follows that p can be written in terms of L and R :

$$p(x) = L(x^2) + xR(x^2) \quad (4)$$

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Recall that we are trying to evaluate p at n roots of unity.

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Recall that we are trying to evaluate p at n roots of unity. Suppose we have a function that takes as input a list of coefficients and returns the evaluation at n roots of unity.

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Recall that we are trying to evaluate p at n roots of unity. Suppose we have a function that takes as input a list of coefficients and returns the evaluation at n roots of unity. We can define this function in terms of itself, because we have a recurrence relation - divide the list in two, giving us L evaluated at $\frac{n}{2}$ th roots of unity and the same for R (from the fact that a n th root of unity squared is a $\frac{n}{2}$ th root of unity).

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Recall that we are trying to evaluate p at n roots of unity. Suppose we have a function that takes as input a list of coefficients and returns the evaluation at n roots of unity. We can define this function in terms of itself, because we have a recurrence relation - divide the list in two, giving us L evaluated at $\frac{n}{2}$ th roots of unity and the same for R (from the fact that a n th root of unity squared is a $\frac{n}{2}$ th root of unity). Finally, we can reconstruct p from L and R according to (4).

Accounting for Edge Cases

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This works directly for ω_n^0 to $\omega_n^{\frac{n}{2}-1}$, however for a power greater than $\frac{n}{2} - 1$ we need to put it in terms of a power less than $\frac{n}{2}$ (since L and R are only $\frac{n}{2}$ long).

Derivation of Negative Property

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Luckily,

$$\begin{aligned}\omega_n^{k+\frac{n}{2}} &= \cos\left(2\pi\frac{k+\frac{n}{2}}{n}\right) + i\sin\left(2\pi\frac{k+\frac{n}{2}}{n}\right) \\ &= \cos\left(\frac{2\pi k}{n} + \pi\right) + i\sin\left(\frac{2\pi k}{n} + \pi\right) \\ &= -\omega_n^k\end{aligned}$$

Putting it All Together

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So, for some power k of the base root of unity we can compute

$$p(\omega_n^k) = L(\omega_n^{2k}) + \omega_n^k R(\omega_n^{2k})$$

and, using the negative property derived before,

$$p(\omega_n^{k+\frac{n}{2}}) = L(\omega_n^{2k}) - \omega_n^k R(\omega_n^{2k})$$

Base Case

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We compute L and R recursively, and we're done!
However, we need to have a base case for the function to work.

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We compute L and R recursively, and we're done!

However, we need to have a base case for the function to work.

Recall our function takes in a coefficient list of length n representing a polynomial, and returns the evaluation of that polynomial at each of the n distinct n th roots of unity.

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We compute L and R recursively, and we're done!

However, we need to have a base case for the function to work.

Recall our function takes in a coefficient list of length n representing a polynomial, and returns the evaluation of that polynomial at each of the n distinct n th roots of unity.

The simplest base case is probably $n = 1$, at which we can stop dividing the list in half and just evaluate the polynomial directly.

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The simplest base case is probably $n = 1$, at which we can stop dividing the list in half and just evaluate the polynomial directly.

The only 1st root of unity is 1, and evaluating a polynomial at $x = 1$ is equal the sum of the coefficients, which for a polynomial with one coefficient is just its singular coefficient.

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We compute L and R recursively, and we're done!

However, we need to have a base case for the function to work.

Recall our function takes in a coefficient list of length n representing a polynomial, and returns the evaluation of that polynomial at each of the n distinct n th roots of unity.

The simplest base case is probably $n = 1$, at which we can stop dividing the list in half and just evaluate the polynomial directly.

The only 1st root of unity is 1, and evaluating a polynomial at $x = 1$ is equal the sum of the coefficients, which for a polynomial with one coefficient is just its singular coefficient.

Base case

Thus, we can just return the coefficient list of the polynomial, or the input to the function when $n = 1$.

Analysis of Runtime

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At every level of the recursion we divide the list in half, making the depth of the recursive tree $\log n$.

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At every level of the recursion we divide the list in half, making the depth of the recursive tree $\log n$.

At a particular level k , we have 2^k nodes and each node has a list of length $\frac{n}{2^k}$, so the total cost of merging the lists together on that level is $2^k \frac{n}{2^k} = n$.

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At every level of the recursion we divide the list in half, making the depth of the recursive tree $\log n$.

At a particular level k , we have 2^k nodes and each node has a list of length $\frac{n}{2^k}$, so the total cost of merging the lists together on that level is $2^k \frac{n}{2^k} = n$.

Thus, $O(\log n \cdot n) = O(n \log n)$.

This algorithm has the same recursive pattern as merge sort.

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So we can evaluate a polynomial at n roots of unity in $O(n \log n)$ with the above algorithm, called the **FFT**.

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So we can evaluate a polynomial at n roots of unity in $O(n \log n)$ with the above algorithm, called the **FFT**.

If we want to multiply two polynomials f and g , we can compute $\text{FFT}(f) \circ \text{FFT}(g)$, where \circ is the element-wise multiplication of the outputs in the point-value representations.

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How do we interpolate coefficients from this point-value representation to complete our convolution?

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How do we interpolate coefficients from this point-value representation to complete our convolution?

We need the inverse FFT, which luckily can be written in terms of the FFT. Recall that the FFT essentially computes the multiplication of the Vandermonde matrix with the coefficients to get to the outputs, e.g. $V\vec{a} = \vec{y}$.

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How do we interpolate coefficients from this point-value representation to complete our convolution?

We need the inverse FFT, which luckily can be written in terms of the FFT. Recall that the FFT essentially computes the multiplication of the Vandermonde matrix with the coefficients to get to the outputs, e.g. $V\vec{a} = \vec{y}$.

To go from the outputs to the coefficients, we can simply multiply by V^{-1} , i.e. $\vec{a} = V^{-1}\vec{y}$.

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Computing V^{-1} is tedious and I don't have much insight (read *Introduction to Algorithms* for a proper proof), but it essentially involves just the definition of matrix inverse and more properties of roots of unity.

Inverse FFT

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Computing V^{-1} is tedious and I don't have much insight (read *Introduction to Algorithms* for a proper proof), but it essentially involves just the definition of matrix inverse and more properties of roots of unity.

It turns out that V^{-1} is essentially V but evaluated at x^{-1} instead of x . Also, divide by n .

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Computing V^{-1} is tedious and I don't have much insight (read *Introduction to Algorithms* for a proper proof), but it essentially involves just the definition of matrix inverse and more properties of roots of unity.

It turns out that V^{-1} is essentially V but evaluated at x^{-1} instead of x . Also, divide by n .

So we can just use the FFT but take the inverse of the root of unity, and divide each element by n at the end.

Inverse FFT

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It turns out that V^{-1} is essentially V but evaluated at x^{-1} instead of x . Also, divide by n .

So we can just use the FFT but take the inverse of the root of unity, and divide each element by n at the end.

Finally, we arrive at the FFT formulation of convolutions.

Convolution Theorem

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Theorem

$f * g = FFT^{-1}(FFT(f) \circ FFT(g))$, i.e. convolutions can be done with FFTs in time $O(n \log n)$.

Proof.

Follows from the presentation up to this point. ☐

A concrete implementation can be found [here](#).

Iterative FFT

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The recursive algorithm can be made iterative surprisingly elegantly from a pattern in binary form of the indexes when recursively subdividing.

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The recursive algorithm can be made iterative surprisingly elegantly from a pattern in binary form of the indexes when recursively subdividing.

I omit the details here, although it makes the algorithm $O(n)$ in memory instead of $O(n \log n)$ and will likely run faster than the recursive algorithm.

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The recursive algorithm can be made iterative surprisingly elegantly from a pattern in binary form of the indexes when recursively subdividing.

I omit the details here, although it makes the algorithm $O(n)$ in memory instead of $O(n \log n)$ and will likely run faster than the recursive algorithm.

An implementation is above.

Number Theoretic Transform (NTT)

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Another improvement on the FFT comes from the observation that complex roots of unity were an arbitrary pick, any **field** with sufficient properties will do.

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Another improvement on the FFT comes from the observation that complex roots of unity were an arbitrary pick, any **field** with sufficient properties will do.

In particular, we can pick a large prime number p and find an equivalent to a root of unity under the field modulo p .

Uses of the NTT

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The details are incredibly tedious and number theory heavy, but they yield the **number theoretic transform**, a variant of the FFT which operates on integers.

Use cases of the NTT

The NTT is useful for polynomials of integer coefficients or certain types of data, e.g. music or images, which have integer pixel values.

Accounting for Negative Numbers

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One downside is that negative numbers do not exist under modulo, which can be accounted for by assuming large numbers are in fact negative, changing the range from $[0, p)$ to $[-\frac{p}{2}, \frac{p}{2})$.

accounting for negative values

```
def ntt_sign(l: list, p: int) -> list:
    return [x if x < (p >> 1) else x - p for x in l]
```

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Before we apply 2D convolutions to images, we elucidate the 1D convolution and its usefulness through an illustrative example.

The anime music quiz problem.

We have a song that is 1 minute and 30 seconds long, and a 10 second clip from that song. We wish to compute:

The Anime Music Quiz Problem

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Before we apply 2D convolutions to images, we elucidate the 1D convolution and its usefulness through an illustrative example.

The anime music quiz problem.

We have a song that is 1 minute and 30 seconds long, and a 10 second clip from that song. We wish to compute:

- 1 Out of a list of songs, which song the clip came from.

The Anime Music Quiz Problem

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Before we apply 2D convolutions to images, we elucidate the 1D convolution and its usefulness through an illustrative example.

The anime music quiz problem.

We have a song that is 1 minute and 30 seconds long, and a 10 second clip from that song. We wish to compute:

- 1 Out of a list of songs, which song the clip came from.
- 2 From a known song, the timestamp where the clip occurred.

Example Run

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```
(amq) stephenhuan@MacBook-Pro ~/P/p/p/amq (master o)> python db.py clip
Picking a clip from NeonGenesisEvangelion at -4.04dB
```

(a) Generating a clip from an anime intro.

NekomonogatariKuro-OP1	: 4590137	loss, occurs at 3.4	seconds in
NeonGenesisEvangelion	: 3931757	loss, occurs at 35.2	seconds in
Nichijou_op2	: 8316782	loss, occurs at 42.4	seconds in
Nisemonogatari_op1	: 4663729	loss, occurs at 77.3	seconds in

```
-----
Final answer: Neon Genesis Evangelion
123 songs in 8.61 seconds = 14.29 songs per second
```

(b) Comparison between songs; finds it occurs exactly 35.2 seconds in.

(c) Song with the lowest loss.

Figure: An example run of the system.

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First, some basics about the representation of audio data.

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First, some basics about the representation of audio data.
We will use the mp3 file format at a sample rate of 48kHz.

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First, some basics about the representation of audio data. We will use the mp3 file format at a sample rate of 48kHz. Audio is fundamentally just a list of numbers, where each number represents the amplitude of the sound wave at that time. A 48kHz sample rate means there are 48,000 of these measurements per second.

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First, some basics about the representation of audio data. We will use the mp3 file format at a sample rate of 48kHz. Audio is fundamentally just a list of numbers, where each number represents the amplitude of the sound wave at that time. A 48kHz sample rate means there are 48,000 of these measurements per second. Each number is a 16-bit number in the range $[0, 1)$, which we will transform to an integer in the range $[0, 2^{16})$ for the NTT.

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So we have two lists of integers, and now wish to find where the smaller list “fits” into the larger list the best.

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So we have two lists of integers, and now wish to find where the smaller list “fits” into the larger list the best.

One way to do this is to compute the ℓ^2 norm, or the vector difference between the two lists.

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So we have two lists of integers, and now wish to find where the smaller list “fits” into the larger list the best.

One way to do this is to compute the ℓ^2 norm, or the vector difference between the two lists.

So we slide the smaller list over the larger list, computing the sum of squares error as we go. Note that this is very similar to the convolution, except we calculate the sum of squares instead of the element-wise product. We also need to flip one of the lists because the convolution flips a list.

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How do we reduce sum of squares to an element-wise product?

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Turning Sum of Squares into Convolutions

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How do we reduce sum of squares to an element-wise product?

We notice that for elements of the lists a, b

$$(a_i - b_j)^2 = a_i^2 - 2a_i b_j + b_j^2$$

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How do we reduce sum of squares to an element-wise product?

We notice that for elements of the lists a, b

$$(a_i - b_j)^2 = a_i^2 - 2a_i b_j + b_j^2$$

When we sum over the length of a , assuming a is the smaller list, we get:

$$\|a\|^2 - 2a \cdot b' + \|b'\|^2$$

where b' is the slice that a overlaps.

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$$(a_i - b_j)^2 = a_i^2 - 2a_i b_j + b_j^2$$

When we sum over the length of a , assuming a is the smaller list, we get:

$$\|a\|^2 - 2a \cdot b' + \|b'\|^2$$

where b' is the slice that a overlaps. $\|a\|$ is a constant, so it can be ignored. Thus, we only need to compute $a \cdot b'$ and $\|b'\|$.

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where b' is the slice that a overlaps. $\|a\|$ is a constant, so it can be ignored. Thus, we only need to compute $a \cdot b'$ and $\|b'\|$.

$a \cdot b'$ directly follows from a convolution and can be read from $a * b^r$, where b^r is the reverse of b .

Turning Sum of Squares into Convolutions

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How do we reduce sum of squares to an element-wise product?

We notice that for elements of the lists a, b

$$(a_i - b_j)^2 = a_i^2 - 2a_i b_j + b_j^2$$

When we sum over the length of a , assuming a is the smaller list, we get:

$$\|a\|^2 - 2a \cdot b' + \|b'\|^2$$

where b' is the slice that a overlaps. $\|a\|$ is a constant, so it can be ignored. Thus, we only need to compute $a \cdot b'$ and $\|b'\|$.

$a \cdot b'$ directly follows from a convolution and can be read from $a * b^r$, where b^r is the reverse of b . Lastly, if we make sure to scan from left to right, then we can compute $\|b'\|$ by storing an intermediate value, and updating it when we slide a an additional index by subtracting out the front of b' , where a left from, and adding the new value that a covers.

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Minimum ℓ^2

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the minimum ℓ^2 between two lists

```
def min_offset(a: list, b: list) -> tuple:
    N, M = len(a), len(b)
    p = fft(a[::-1], b)[N - 1:]
    x2, xy, y2 = sum(x*x for x in a), p[0], \
                 sum(b[i]*b[i] for i in range(N))
    best, l2 = 0, -2*xy + y2
    for i in range(1, M - N + 1):
        y2 += b[N - 1 + i]*b[N - 1 + i] - b[i - 1]*b[i - 1]
        xy = p[i]
        d = -2*xy + y2
        if d < l2:
            best, l2 = i, d
    return best, x2 + l2
```

NTT Considerations

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We need to be careful about a few things. If we don't pick p for the NTT large enough, then it won't work.

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We need to be careful about a few things. If we don't pick p for the NTT large enough, then it won't work.

If m is the largest number in a list and n is the length of the list, then we need p to be bigger than m^2n , the largest a single element can become.

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We need to be careful about a few things. If we don't pick p for the NTT large enough, then it won't work.

If m is the largest number in a list and n is the length of the list, then we need p to be bigger than m^2n , the largest a single element can become.

n is $90 \cdot 48,000 \approx 4 \cdot 10^6$ and m is 2^{16} .

$m^2n = 2^{32} \cdot 4 \cdot 10^6 \approx 2^{54}$.

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n is $90 \cdot 48,000 \approx 4 \cdot 10^6$ and m is 2^{16} .

$m^2n = 2^{32} \cdot 4 \cdot 10^6 \approx 2^{54}$.

This seems fine since 2^{54} will fit in a long, but this won't work since we need to compute x^2 as part of the FFT, and $(2^{54})^2$ will definitely overflow.

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n is $90 \cdot 48,000 \approx 4 \cdot 10^6$ and m is 2^{16} .

$m^2n = 2^{32} \cdot 4 \cdot 10^6 \approx 2^{54}$.

This seems fine since 2^{54} will fit in a long, but this won't work since we need to compute x^2 as part of the FFT, and $(2^{54})^2$ will definitely overflow.

We could get around this overflow by doing modulo multiplication instead of standard multiplication, but that would introduce a log factor, making the algorithm 64x slower, an unacceptable slowdown.

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One trick is to reduce the bitrate of the mp3 at the sacrifice of audio quality, and go from 16-bit audio to 8-bit audio.

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One trick is to reduce the bitrate of the mp3 at the sacrifice of audio quality, and go from 16-bit audio to 8-bit audio.

A naive way to do it would be to multiply the real number by 2^8 and round, but a better way is the μ -law algorithm, a trick that preserves frequencies closer to the human voice. A comparison between naive scaling and the μ law is presented [here](#).

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A naive way to do it would be to multiply the real number by 2^8 and round, but a better way is the μ -law algorithm, a trick that preserves frequencies closer to the human voice. A comparison between naive scaling and the μ law is presented [here](#).

With 8-bit audio, $m^2 n = 2^{16} \cdot 4 \cdot 10^6 \approx 2^{38}$. This goes over the limit of 2^{32} for x^2 to fit in a long, but it works in practice since mp3 audio rarely hits maximum volume and our clip is 10 seconds long, and we computed for 90 seconds.

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A naive way to do it would be to multiply the real number by 2^8 and round, but a better way is the μ -law algorithm, a trick that preserves frequencies closer to the human voice. A comparison between naive scaling and the μ law is presented [here](#).

With 8-bit audio, $m^2 n = 2^{16} \cdot 4 \cdot 10^6 \approx 2^{38}$. This goes over the limit of 2^{32} for x^2 to fit in a long, but it works in practice since mp3 audio rarely hits maximum volume and our clip is 10 seconds long, and we computed for 90 seconds.

An implementation is given [here](#) and a video walkthrough [here](#).

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2D convolutions, a convolution generalized to matrices, are useful in computer vision for a variety of reasons, including edge detection and convolutional neural networks. Their exact usage will not be discussed here, and instead we will discuss an efficient way to calculate a 2D convolution with the FFT we have already developed.

Definitions

We have an “data” matrix, representing an image, and we have a *kernel* matrix, which is the matrix we imagine sliding over the image. This is also known as a *filter*.

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For 2D convolutions, the result is slightly ambiguous depending on how one defines it.

We will use [scipy's](#) definition, where to calculate the value of the convolution at a particular point, we imagine the *bottom right* corner of the kernel placed over that point.

Convolution Example

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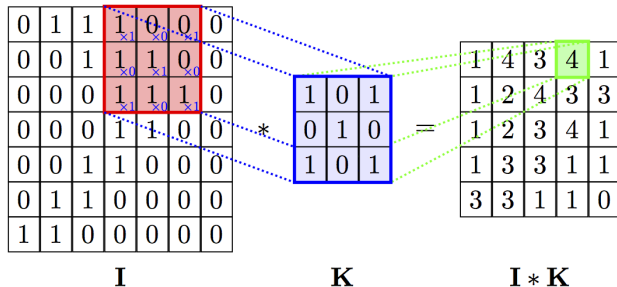


Figure: A convolution taken from [here](#).

Definition of a 2D Convolution

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We define the 2D convolution between an image x of size $M \times N$ and a kernel h of size $H \times W$ as follows (similar to the 1D case, we assume both matrices are padded with 0's):

Definition of the 2D convolution

$$(x * h)[i, j] = \sum_{k=0}^i \sum_{l=0}^j x[k][l] h[i - k][j - l]$$

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This operation is also symmetric, so what we call the image and the kernel is essentially arbitrary (by convention, the kernel is the smaller matrix). The resulting matrix is going to be of size $(M + H - 1) \times (N + W - 1)$ from the same logic as the 1D case.

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This operation is also symmetric, so what we call the image and the kernel is essentially arbitrary (by convention, the kernel is the smaller matrix). The resulting matrix is going to be of size $(M + H - 1) \times (N + W - 1)$ from the same logic as the 1D case. Thus, the time it takes to compute the convolution is $O(MNHW)$. We can, however, take advantage of a trick if the kernel has a certain property.

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Separable

A matrix M is *separable* if it can be written as $\vec{u}\vec{v}^T$ for some vectors \vec{u}, \vec{v} .

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Separable

A matrix M is *separable* if it can be written as $\vec{u}\vec{v}^T$ for some vectors \vec{u}, \vec{v} .

Sobel matrix

The famous Sobel matrix for edge detection is separable:

$$\begin{bmatrix} 1 & 0 & -1 \\ 2 & 0 & -2 \\ 1 & 0 & -1 \end{bmatrix} = \begin{bmatrix} 1 \\ 2 \\ 1 \end{bmatrix} \begin{bmatrix} 1 & 0 & -1 \end{bmatrix}$$

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Separating the convolution

If M is separable, we can convolute with \vec{u} and then with \vec{v} .

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Theorem

*If $h = \vec{u}\vec{v}^T$, then $(x * h) = ((x * \vec{u}) * \vec{v})$, i.e. we can separate a convolution into two parts.*

Proof.

$$(x * u)[i, j] = \sum_{k=0}^i \sum_{l=0}^j x[k][l] u[i - k][j - l] \quad \text{Definition}$$

Since u is a column vector, it only has values when $l = j$, removing the inner sum.

$$= \sum_{k=0}^i x[k][j] u[i - k][0]$$

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Proof.

Convoluting with v ,

$$((x * u) * v)[i, j] = \sum_{k=0}^i \sum_{l=0}^j \left(\sum_{m=0}^k x[m][l] u[k-m][0] \right) v[i-k][j-l]$$

Since v is a row vector, it only has values when $k = i$, removing the outermost sum.

$$= \sum_{l=0}^j \left(\sum_{m=0}^i x[m][l] u[i-m][0] \right) v[0][j-l]$$

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Proof.

Swapping the order of summations and renaming m to k ,

$$= \sum_{k=0}^i \sum_{l=0}^j x[k][l] u[i-k][0] v[0][j-l]$$

From the fact that $h[x][y] = u[x][0] v[0][y]$,

$$\begin{aligned} &= \sum_{k=0}^i \sum_{l=0}^j x[k][l] h[i-k][j-l] \\ &= (x * h) \end{aligned}$$



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How does this help us?

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How does this help us?

Well, recall the running time of $O(MNHW)$. If we do two convolutions of a kernel of $H \times 1$ and another of $1 \times W$, the running time will be $O(MNH + MNW) = O(MN(H + W))$, a significant improvement as HW grows quadratically while $H + W$ grows linearly.

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How does this help us?

Well, recall the running time of $O(MNHW)$. If we do two convolutions of a kernel of $H \times 1$ and another of $1 \times W$, the running time will be $O(MNH + MNW) = O(MN(H + W))$, a significant improvement as HW grows quadratically while $H + W$ grows linearly.

We can also use repeated 1D convolution to compute the 2D convolution for the specific case of a vector, yielding a $O(MN \log MN)$ time algorithm.

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However, not every matrix is separable. The conditions are quite strict, a matrix is separable if and only if every pair of rows is a multiple of each other, i.e. the matrix is made up of multiples of a particular row vector. As a consequence, the matrix is also made up of multiples of a particular column vector. These matrices are relatively rare, so there is utility in deriving a more general algorithm.

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We can reduce 2D convolutions to 1D convolutions if we're clever. The observation is that if we **flatten** both matrices into a 1D list by reading from top to bottom, left to right, we can just convolute in 1D and reconstruct the matrix afterwards.

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We can reduce 2D convolutions to 1D convolutions if we're clever. The observation is that if we **flatten** both matrices into a 1D list by reading from top to bottom, left to right, we can just convolute in 1D and reconstruct the matrix afterwards. We need to make sure both matrices are sufficiently padded with zeros, such that the zeros force values in the kernel to their proper rows in the image.

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We can reduce 2D convolutions to 1D convolutions if we're clever. The observation is that if we **flatten** both matrices into a 1D list by reading from top to bottom, left to right, we can just convolute in 1D and reconstruct the matrix afterwards.

We need to make sure both matrices are sufficiently padded with zeros, such that the zeros force values in the kernel to their proper rows in the image.

It turns out that we can just pad both matrices to the final column size of the convolution, $N + W - 1$, flatten both, convolute with the FFT, and then reshape the resulting list to a matrix of proper size.

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- 1 Pad the rows of both matrices with zeros such that each has a width of $N + W - 1$

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- 1 Pad the rows of both matrices with zeros such that each has a width of $N + W - 1$
- 2 Flatten both by reading top to bottom, left to right

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- 1 Pad the rows of both matrices with zeros such that each has a width of $N + W - 1$
- 2 Flatten both by reading top to bottom, left to right
- 3 Convolute the resulting lists in 1D

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- 1 Pad the rows of both matrices with zeros such that each has a width of $N + W - 1$
- 2 Flatten both by reading top to bottom, left to right
- 3 Convolute the resulting lists in 1D
- 4 Reconstruct a 2D matrix, because we know its shape is $(M + H - 1) \times (N + W - 1)$

The 2D Convolution Algorithm

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```
_____ the 2D convolution algorithm _____  
def flatten(m: list, pad=0) -> list:  
    """ Flattens a matrix into a list. """  
    return [x for row in m for x in row + [0]*pad]  
  
def reshape(l: list, m: int, n: int) -> list:  
    """ Shapes a list into a MxN matrix. """  
    return [[l[r*n + c] for c in range(n)]  
            for r in range(m)]  
  
def conv(h: list, x: list):  
    """ Computes the 2D convolution. """  
    M, N, H, W = len(x), len(x[0]), len(h), len(h[0])  
    # need to pad the columns to the final size  
    h, x = flatten(h, N - 1), flatten(x, W - 1)  
    return reshape(fft(h, x), M + H - 1, N + W - 1)
```

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In most computer vision applications, the kernel is a square matrix of size $K \times K$, where K is an odd number.

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In most computer vision applications, the kernel is a square matrix of size $K \times K$, where K is an odd number.

The *middle value* of the kernel is then placed over each pixel of the image, yielding a transformed image of the same dimensionality as the original.

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In most computer vision applications, the kernel is a square matrix of size $K \times K$, where K is an odd number.

The *middle value* of the kernel is then placed over each pixel of the image, yielding a transformed image of the same dimensionality as the original.

We can simulate this by simply cutting off the first and last $\frac{K-1}{2}$ rows and the same for the columns. This transforms the resulting size from $N + K - 1$ to $N + K - 1 - 2\frac{K-1}{2} = N$.

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pruning

```
def prune(h: list, x: list) -> list:
    """ Prunes a convolution for the specific  $K \times K$ 
        ↪ filter case. """
    m, k = conv(h, x), min(len(h), len(x))
    pad = (k - 1) >> 1
    return [row[pad:-pad] for row in m[pad:-pad]]
```

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The running time of the algorithm is going to be $O(MN \log MN) = O(MN(\log N + \log M)) = O(MN \log N)$ since we convolute a list of length $M(N + W - 1)$, and we assume $N \geq M > W$.

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This is not necessarily faster than the brute-force algorithm; it depends on the kernel size. For simplicity, suppose we have a $N \times N$ image and a $K \times K$ kernel where $N > K$. Brute force yields $O(N^2 K^2)$ while the FFT algorithm yields $O(N^2 \log N)$.

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Thus, if $\log N < K^2$ then the FFT algorithm is going to be faster. For $K > 5$ that is a fair assumption since $K^2 = 25$, 2^{25} is several million. Obviously the FFT algorithm has a much larger constant factor, but for a sufficiently large kernel the time savings become greater and greater.

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The convolution, an operator very useful for signal, audio, and image processing, can be efficiently computed with the Fast Fourier Transform, or FFT.

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The convolution, an operator very useful for signal, audio, and image processing, can be efficiently computed with the Fast Fourier Transform, or FFT.

If the data is integer, then floating-point arithmetic can be avoided with the Number Theoretic Transform (NTT), a variant of the FFT which uses modulo instead of complex numbers, and calculates entirely in integers.

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This lecture skips over the continuous case (what I've been calling the Fast Fourier Transform is more mathematically called the **Discrete Fourier Transform**, or DFT) but the idea is essentially the same, any summation turns into an integral.

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This lecture skips over the continuous case (what I've been calling the Fast Fourier Transform is more mathematically called the **Discrete Fourier Transform**, or DFT) but the idea is essentially the same, any summation turns into an integral. It also skips over the mathematical interpretation of the FFT, involving decomposing a function into a series of sine and cosine waves. This is useful for signal processing and audio analysis, but requires a stronger mathematical background and to be honest, I haven't studied it at all myself. Fourier analysis goes deeper than we need here.

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Introduction to Algorithms is definitely the most helpful source on the FFT (from a computer science perspective), and more thorough treatments of the FFT from an engineering or mathematical standpoint are not hard to find.

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1 [SPOJ POLYMUL](#): Direct application of the FFT.

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- 2 **SPOJ MUL**: Given 1000 pairs of numbers, compute the product of each pair; each number can have up to 10,000 digits.

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- 2** Solution: Think of numbers as polynomials, where the digits are coefficients and x is 10. Then, you can multiply two numbers by multiplying the polynomials. However, there is no guarantee that the coefficients of the resulting polynomial are less than 10, so it is not a valid number. As a last post-processing step, start from the smallest place value and move your way to the largest, moving the digit overflow from one place value to the next. Since you iterate over the number of digits in the number, it takes $O(\log n)$ which is dominated by the FFT.

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- 2 An extension of this idea is the [Schönhage–Strassen](#) algorithm, which disregards the requirement that the intermediate numbers fit in a long, at the cost of being $O(n \log n \log \log n)$. A more recent algorithm, by Harvey and van der Hoeven, achieves $O(n \log n)$.

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- 3 **SPOJ MAXMATCH**: Given a string S of length N made up of the characters “a”, “b”, and “c”, compute the maximum self-matching, where a self-matching is defined as the number of characters which match between S and S shifted some nonzero number of characters.

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- 3** Solution: For an offset i , the size of the overlap will be $N - i$. So we just need to find the number of differences, and subtract that from $N - i$ to obtain the number of matches. The easiest thing to do is to keep track of each character separately, so to compute the differences for each character. Suppose our character is “a”. We encode “a” as a 0, and the other characters as a 1. We then find the ℓ^2 norm between this new list and this list with N 1’s added to it (so that when we overlap, the non “a” characters aren’t counted). This has the complication of counting “a”’s which are off the edge of the string, which we can account for by simply keeping track of the number of “a”’s we have seen.

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- 3 Given $a[i]$ as the number of mismatches with the character “a” at a shift of i , and $b[i], c[i]$, the number of matches is $N - i - \frac{a[i] + b[i] + c[i]}{2}$. We divide by 2 because we count each mismatch twice (once for each character in the pair).

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- 3** A much conceptually simpler algorithm is to encode “a”, “b”, and “c” cleverly and then compute the matches in one shot. If we encode “a” as $(1, 0, 0)$, “b” as $(0, 1, 0)$, and “c” as $(0, 0, 1)$, the FFT of the resulting list with its reverse will give us the number of matches at each index because the character representations dot each other will be 1 if they are equal, and 0 if they are unequal. Thus, the FFT will give us exactly the number of matches, but we need to only look at every 3rd index since the other 2 are byproducts of our transformation.
- In practice, running one big FFT is faster than running 3 smaller FFTs.

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- 4 [Codechef FARASA](#): Given an array, find the number of distinct sums of a contiguous subarray.

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4 Solution: [editorial](#).

Fair warning, time bounds are ridiculous.

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- 5 [Codeforces Round #296](#): Given two strings T, S and an error bound k , find all the positions where T occurs in S , where T “occurring” at some index i means that the j th character of T has a corresponding character within k of its position.

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5 Solution: Honestly no clue but it has the “FFT” tag.

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- 6 **String matching with wildcards:** Given two binary strings T, S , T has length N and has wildcards which match any character in S , find all occurrences of T in S .

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- 6** Solution: Encode 1 as 1 and 0 as -1. The dot product between T and the slice that T overlaps with S will be N if they match exactly and less than N if they don't match exactly. To account for wildcards, encode a wildcard as 0 and count the number of wildcards, C . Then, if they match exactly it will be $N - C$, and less than that if they don't.

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- 6** This can be generalized to non-binary strings if you apply the above algorithm to each character, setting that character as 1 and not that character as -1. Sum over all possible characters, and that will tell you whether there is a mismatch somewhere (similar to SPOJ MAXMATCH).

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- 6 This idea can also be applied to string matching without wildcards. Encode each character as its ASCII value in a polynomial, and compute the ℓ^2 -norm between T and S . The ℓ^2 norm will be 0 if they match, and positive if they don't.

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- 7 **3SUM**: Given a list of integers between $-N$ and N , find 3 numbers that add up to 0 (duplicates are allowed).

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- 7 Solution: The basic idea will be to encode the list into a length $2N$ polynomial p where the degree is an integer value and the coefficient is whether that value appears in the array. Compute p^3 and read off the coefficient of x^0 .

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- 7 However, this doesn't work if the degrees are negative. If the most negative power of x in p is x^{-N} , We can simply multiply p by x^N to make every power positive, making a new polynomial p' . Then, after computing $(p')^3$, instead of looking at the coefficient of x^0 , we can look at the coefficient of x^{-3N} (accounting for the fact that $p' = x^N p$, $(p')^3 = x^{3N} p^3$, $p^3 = \frac{(p')^3}{x^{3N}}$)

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- 7** Alternative solution, if duplicates aren't allowed: [here](#) (look for “color coding”).

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- 8 **Anime Music Quiz:** Guess which anime an intro/outro comes from.

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8 Solution: The [Shazam algorithm](#).

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- 1 “Edge Detection”, (Alexey Didenkov, 2018)
- 2 “Fast Multiplication: Karatsuba and FFT” (Haoyuan Sun, 2016)
- 3 “Multiplying Polynomials”, (Haoyuan Sun, 2015)
- 4 “Fast Fourier Transform”, (Sreenath Are, 2013)

References

Resources that were useful when compiling this lecture

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- 1 *Introduction to Algorithms*, chapter 30 (very helpful)
- 2 The number theoretic transform
- 3 μ -law algorithm
- 4 Picard's Existence and Uniqueness Theorem
- 5 Separable convolutions