

7.3

February 21, 2019

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In [1]: # Content from Proakis
        # Code l 2019, Alexander Kain
        import numpy as np
        from numpy.fft import fft, ifft, rfft, irfft

        from matplotlib import pyplot as plt
        %matplotlib inline
        plt.rcParams['figure.figsize'] = (10.0, 8.0)

        import sympy as sym
        sym.init_printing(use_unicode=True)
```

0.1 7.3 Linear Filtering Methods Based on the DFT

We can use the DFT as a computational tool for linear filtering of systems with finite impulse responses. Due to the existence of the FFT this approach is often more efficient than time-domain convolution.

0.1.1 7.3.1. Use of the DFT in Linear Filtering

We know that the product of two DFTs is equivalent to circular convolution. We seek a methodology that is equivalent to linear convolution.

Suppose we have $x[n]$ of length L which excites an FIR filter with impulse response $h[n]$ of length M :

$$x[n] = 0, \quad n < 0 \text{ and } n \geq L$$

$$h[n] = 0, \quad n < 0 \text{ and } n \geq M$$

In time domain, the output sequence $y[n]$ can be computed by convolving $x[n]$ and $h[n]$

$$y[n] = \sum_{k=0}^{M-1} h[k]x[n-k]$$

The length of $y[n]$ will be exactly $L + M - 1$.

The frequency domain equivalent is

$$Y(\omega) = X(\omega)H(\omega)$$

If the sequence $y[n]$ is to be represented uniquely in the frequency domain by samples of its spectrum $Y(\omega)$, the number of distinct samples must equal or exceed $L + M - 1$. Therefore, a DFT of size $N \geq L + M - 1$ is required and

$$Y[k] = X[k]H[k]$$

where $X[k]$ and $H[k]$ are the N -point DFTs of the sequences $x[n]$ and $h[n]$.

This implies that the $N \geq L + M - 1$ -point circular convolution of $x[n]$ with $h[n]$ is *equivalent* to linear convolution.

In [2]: # *Example 7.3.1*

```
x = np.r_[1, 2, 2, 1]
h = np.r_[1, 2, 3]
L = len(x)
M = len(h)
N = L + M - 1

# linear convolution
y = np.zeros(N)
for n in range(N):
    for k in range(M):
        ix = n-k
        if 0 <= ix < L:
            y[n] += h[k] * x[ix]

assert np.allclose(y, np.convolve(x, h)) # TD library implementation
y
```

Out[2]: array([1., 4., 9., 11., 8., 3.])

In [3]: # *aside: list comprehension version*

```
y = np.array([
    np.sum([h[k] * x[n-k] for k in range(M) if 0 <= (n-k) < L])
    for n in range(N)])

assert np.allclose(y, np.convolve(x, h))
```

In [4]: $X = \text{rfft}(x, N)$ # *equivalent to zero-padding*

$H = \text{rfft}(h, N)$

$Y = X * H$

$y = \text{irfft}(Y)$

```
from scipy.signal import fftconvolve
assert np.allclose(y, fftconvolve(x,h)) # FD library implementation
y
```

Out[4]: array([1., 4., 9., 11., 8., 3.])

```

In [5]: fftn = 1 << (N-1).bit_length() # FFT performs best with powers of two
        print(fftn)

        X = rfft(x, fftn)
        H = rfft(h, fftn)
        Y = X * H
        y = irfft(Y)[:N] # keep only what is needed

        assert np.allclose(y, fftconvolve(x,h)) # FD library implementation
        y

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Out[5]: array([ 1.,  4.,  9., 11.,  8.,  3.])

```

0.1.2 7.3.2. Filtering of long Data Sequences

A long signal must be segmented to fixed-size blocks prior to processing, which will then be processed one *segment/block/frame* at a time, and then fit back together.

Overlap-add method In this method, the size of the block is L (it is assumed that $L \gg M$), and the size of the DFTs is $N \geq L + M - 1$.

The *last* $M - 1$ points from each block must be overlapped and added to the *first* $M - 1$ points of the succeeding block.

