Assignment 10: Convolution

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1 Aim

The goal is to implement linear and circular convolution using methods of scientific computing. Convolution is defined as

$$y[n] = \sum_{k=-\infty}^{\infty} x[n-k]h[k]$$

In Fourier domain,

$$Y[m] = X[m]H[m]$$

2 The Assignment

2.1 The FIR Filter

The h.csv file is given on Moodle. It contains the FIR h[n]. Then I take a DFT using the sig.freqz() function. The magnitude and phases are plotted (Figure 1 and 2). The code follows.

```
with open("h.csv") as f:
    h = np.loadtxt(f, delimiter=",")

# plotting the filter response
# USE: scipy.signal.freqz()
w, H = sig.freqz(h) # takes the fourrier transform of the filter
plt.plot(w, 20 * np.log10(abs(H)))
plt.xlabel("Frequency (rad/sample)")
plt.ylabel("Magnitude (dB)")
plt.title("Magnitude Response")
plt.grid()
plt.savefig("h.png")
plt.show()
```

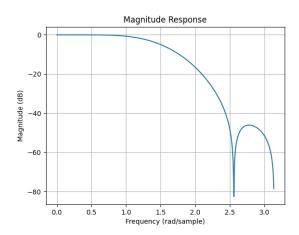


Figure 1: Magnitude Response

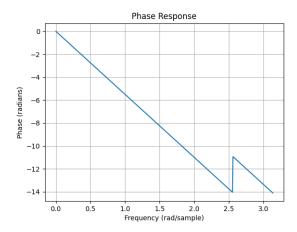


Figure 2: Phase Response

2.2 Input Signal

$$x[n] = \cos(0.2\pi n) + \cos(0.85\pi n)$$

n varies from 1 to 2^{10} . The sequence is generated and plotted.

2.3 Linear Convolution

The output of the LTI system is the convolution of input and impulse response. A computationally intensive idea is nested for loops.

```
y = np.zeros(len(x))
for i in range(len(x)):
    for j in range(len(h)):
```

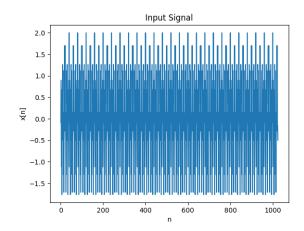


Figure 3: Input Signal

The high frequency (0.85) is filtered out.

2.4 Circular Convolution

I zero pad the FIR such that x and h are now equally long. Then their DFTs are multiplied, IDFT gives the required output. It looks the same as the one

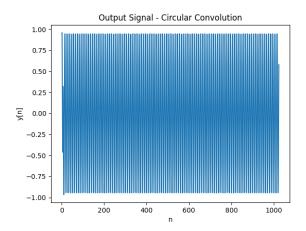


Figure 4: Output Signal

each section is 16 2.5 Linear Convolution Using Circular Convolution samples, L = 64

This is tricky. Our FIR was 12 samples long (P = 12). To extend it to 2^m , we need to add 4 more zeros. Hence it is now 16 samples. This is h_with_zeros. Now, the input signal had 1024 samples. I make sections of x such that each is L = 64 long. In our case, x had 2^m samples already, which may not be the case always. Hence, the line to zero pad. Now begins the circular convolution algorithm. The logic is explained in the assignment how y will have N wrong values and how it obtains the correct numbers from the next computation. I take real values of the output for the plot's sake and plot the first 1024 values of y_circ_conv (as n is defined that way). Note that the length of y_circ_conv is 1039, which is len(x) + len(h) - 1.

each succeeding convolution adds to the previous values of y.

```
(x_{int} + 1) * len_h : (i + 1) * len_h],
         np.zeros(len_h - 1))
    y\_circ\_conv[i * len\_h : (i + 1) * len\_h + len\_h - 1] +=
     np.fft.ifft(
        np.fft.fft(x_)
        * np.fft.fft(
            np.concatenate((h_with_zeros, np.zeros(len(x_))
             - len(h_with_zeros))))
        )
    ).real
11 11 11
if .real is not used, the following error is thrown:
    numpy.core._exceptions.UFuncTypeError: Cannot cast
     ufunc 'add' output from dtype('complex128') to
     dtype('float64') with casting rule 'same_kind'
11 11 11
plt.plot(n, (y_circ_conv[:1024]).real)
                                          \# as n goes from 1 to 2**10
plt.xlabel("n")
plt.ylabel("y[n]")
plt.title("Linear Convolution from Circular Convolution")
plt.savefig("lin_from_cir.png")
plt.grid()
plt.show()
```

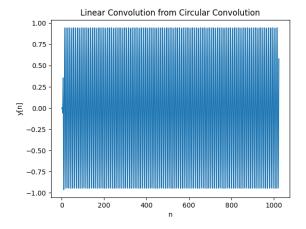


Figure 5: Liinear from Circular Convolution

2.6 Circular Correlation

Open the Zadoff - Chu sequence file, obtain the complex numbers from that.

```
with open("x1.csv") as f:
    csvreader = csv.reader(f)
    lines = np.array([row for row in csvreader])
# print(lines, type(lines), lines.shape)
# this tells that the array is a numpy array
  and numbers are given as a+ib
# we need to convert them to complex numbers
zandoff_chu_seq = []
for line in lines:
    try:
        a, bi = line[0].split("+")
        a = float(a)
        bi = float(bi[:-1]) # leaving the i
    except ValueError:
        try:
            if line[0][0] == "-":
                _, a, bi = line[0].split("-")
            else:
                a, bi = line[0].split("-")
            a = float(a)
            bi = -float(bi[:-1]) # leaving the i
        except ValueError:
            a = float(line[0]) # there is a 1 in the list
            bi = 0.0
    zandoff_chu_seq.append([a + 1j * bi])
  Now, compute correlation.
zandoff_chu_seq = np.array(zandoff_chu_seq)
# print(zandoff_chu_seq, len(zandoff_chu_seq), zandoff_chu_seq.shape)
# correlation of this sequence with a shifted version of itself
shifted_zandoff_chu_seq = np.roll(zandoff_chu_seq, 5)
correlation = np.fft.ifftshift(
    np.correlate(shifted_zandoff_chu_seq[:, 0],
    zandoff_chu_seq[:, 0], "full")
)
print(len(correlation))
plt.plot(np.arange(0, len(correlation)), abs(correlation), "ro")
```

```
plt.xlabel("n")
plt.ylabel("correlation")
plt.xlim(0, 30)
plt.title("Correlation of Zadoff-Chu Sequence")
plt.savefig("correlation.png")
plt.grid()
plt.show()
```

As expected, it gives a peak at 5.

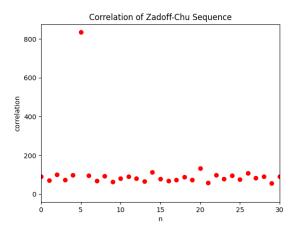


Figure 6: Correlation

3 Conclusion

Direct computation of linear convolution has a high time complexity. A faster way to do this is using DFTs. Also, circular convolution can be used to compute the output at a much lower (log(n)) complexity. Towards the end, an interesting property of the Zadoff Chu sequence was studied that it shows a peak in its auto-correlation at the point corresponding to the shift.