# Write a code for correlation of two given DTSs without using inbuilt function.

$$y(n)=x(n)*h(-n)=\sum_{k=-\infty}^{\infty}x(k)*h(n-k)$$

## Given data

```
Discrete time sequences 1 (DTS1): X_1(n_1) = \{1,2,3,4\} , n_1=-1:2
```

Discrete time sequences 2 (DTS2):  $X_2(n_2)$  =  $\{1,2,1,1\}$  ,  $n_2=-2:1$ 

Before solving it, let me explain my approach. As per the formula, Correlation is same as the convolution of X(n) and H(-n), I will pass these values to our prevously done convolution function.

# **Import Modules**

```
In [1]:
```

```
import numpy as np
import matplotlib.pyplot as plt
```

# Helper function for plotting

```
In [2]:
```

# **Folding**

#### In [20]:

```
def folding(arr, n):
    # sanity check for length of timestamp and signal array must be same
    assert len(arr) == len(n)
    n = np.array(n)
    index_0 = np.where(n == 0)[0][0]
    # initiate an empty array to store our updated array
    new arr = []
    # push all the positive timestamps to our new arr
    # for example in our case
    # [-1, 0, 1, 2] will be [-2, -1, 0, 1] and corresponding array values will be updat
ed accordingly
    # push from 2 to 1 in reverse order
    for i in range(len(n) - 1, index_0 - 1, -1):
        new_arr.append(arr[i])
    # push from 0 to -1 in the same order
    for j in arr[:index_0][::-1]:
        new arr.append(j)
   # our n is now [2, 1, 0, -1] hence return the negative sorted values i.e [-2, -1, -1]
 0, 1] which is our folded timestamp
    return np.array(new_arr), sorted(-n)
```

## **Padding**

#### In [4]:

#### **Calculate the Convolution**

#### In [5]:

```
def calculate_convolution(x, h):
   parameters:
   x -> zero padded signal of len = max_len i.e l1 + l2 - 1
    h -> our folded h(n) signal
    # initiate the matrix as said above
    ans = np.zeros((x.shape[0], x.shape[0]))
    # keep stacking the linear kernels which will be later passed on our padded signal
    for i in range(ans.shape[0]):
        if i < len(h):</pre>
            ans[i, :i + 1] = h[-i - 1:]
        else:
            ans[i, i - len(h)+1: i+1] = h
    # print to observe
    print('\nStack of linear kernels: \n\n', ans)
    # take the transpose
    ans = np.transpose(ans)
    # return matrix multiplication of our padded signal and our kernel matrix, this is
 our desired result
    return np.matmul(x, ans)
```

# Putting it all together

#### In [7]:

```
def main(x1, n1, x2, n2):
    Calculates the convolution of x1 and x2 where
   x1 -> original signal
    x2 -> filter/kernel
    n1 -> timestamps of x1
    n2 -> timestamps of x2
    # show signal x1
    print("Signal x is: ", x1)
    print("\nTime stamp of X is: ", n1)
    plot_graph(x1, n1, y_label='$x1$', x_label='timestamp', graph_title='Signal $x$')
    # show signal x2
    print("\nSignal H is: ", x2)
    print("\nTimestamp of H is: ", n2)
    plot_graph(x2, n2, y_label='$h(n)$', x_label='timestamp', graph_title='Signal $h$')
    # apply folding on second signal
    folded_H, neg_timestamp = folding(x2, n2)
    print("\nFolded H is: ", folded_H)
    print("\nTimestamp of H is: ", neg_timestamp)
    plot_graph(folded_H, neg_timestamp, x_label='timestamp', y_label='$h(-n)$', graph_t
itle='Folded $h(n)$')
    # create the resutant timestamp
    # take the min of min of timestamps
    left_most_timestamp = min(min(n1), min(neg_timestamp))
    # calculate the maximum sequence length
   max_len_result = len(n1) + len(neg_timestamp) - 1
    # initialize the resultant convolution array with zeroes
    result_time_stamp = np.arange(left_most_timestamp, left_most_timestamp + max_len_re
sult, 1)
    print("\nResultant timestamp would be: ", result_time_stamp)
    # pad our Signal X upto the len of max_len
    # 3 is obtained from result time stamp - len(x1)
   X = zero pad(x1, right pad=3)
    # Here is our result
    result = calculate_convolution(X, folded_H)
    print('\nConvolution of Signal x1 and x2 is: ', result)
    plot graph(result, result time stamp, x label='timestamp', y label='$\sum x(k) * h
(n-k)$', graph title='Convoution Result')
    assert all(np.convolve(x1, x2) == result), 'Computed Result is Wrong'
```

As our main function calculates the convolution of two signals, use it to calculate the convolution of X(n) and H(-n), which is our desired result i.e correlation of X(n) and H(-n).

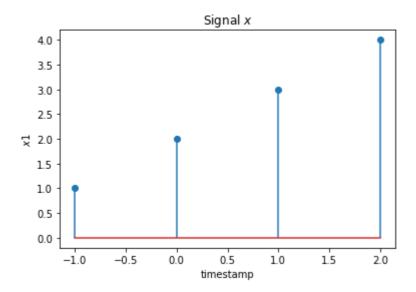
#### In [8]:

```
def correlation_main(x, n1, h, n2):
    Calculates the corelation of x and h where
       x -> original signal
       h -> filter/kernel
       n1 -> timestamps of x
       n2 -> timestamps of h
    steps to follow:
        1. flip our signal H(n) given to get H(-n)
        2. Call our main() function to calculate the convolution of the original signal
and the flipped signal
    ###################
    # Flip the signal H(n) to get H(-n)
   flipped_h, flipped_n = folding(h, n2)
    # call our convolution function that we defined above
    # pass the arguments, x as main signal and flipped_h as kernel
   main(x, n1, flipped_h, flipped_n)
    pass
```

#### In [21]:

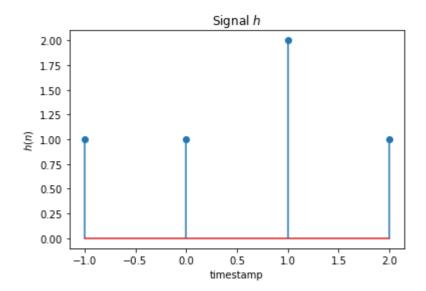
Signal x is: [1 2 3 4]

Time stamp of X is: [-1 0 1 2]



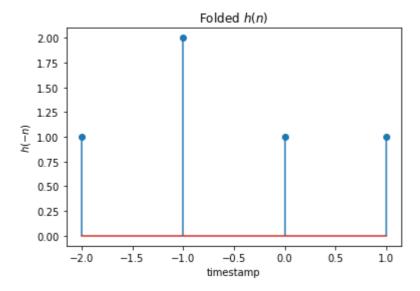
Signal H is: [1 1 2 1]

Timestamp of H is: [-1, 0, 1, 2]



Folded H is: [1 2 1 1]

Timestamp of H is: [-2, -1, 0, 1]

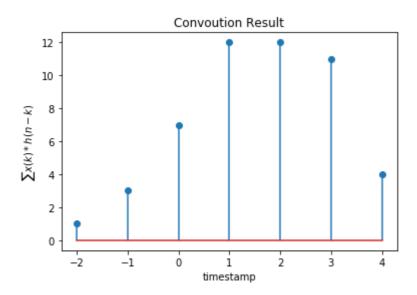


Resultant timestamp would be: [-2 -1 0 1 2 3 4]

Stack of linear kernels:

[[1. 0. 0. 0. 0. 0. 0. 0.] [1. 1. 0. 0. 0. 0. 0. 0.] [2. 1. 1. 0. 0. 0. 0.] [1. 2. 1. 1. 0. 0. 0.] [0. 1. 2. 1. 1. 0. 0.] [0. 0. 1. 2. 1. 1. 0.] [0. 0. 0. 1. 2. 1. 1.]

Convolution of Signal x1 and x2 is: [ 1. 3. 7. 12. 12. 11. 4.]



# Kindly read the comments and docstring of the functions.

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