## ps6

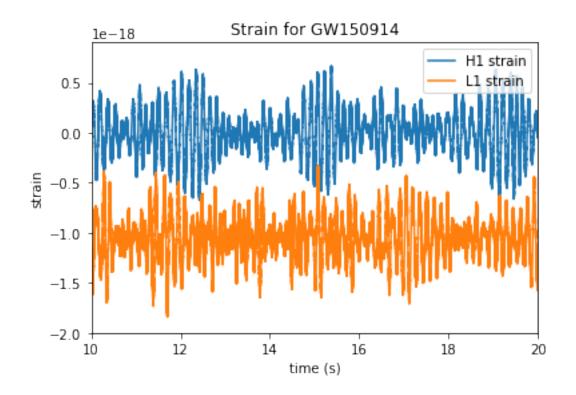
## November 5, 2021

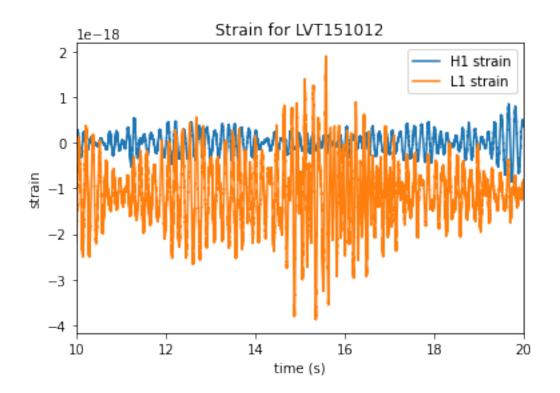
```
[1]: import numpy as np
     import matplotlib.pyplot as plt
     import h5py
     from scipy.signal import savgol_filter
     import json
     import glob
     datadir = 'LOSC_Event_tutorial/'
[2]: # Copied from the class example
     def read_template(filename):
         dataFile = h5py.File(filename, 'r')
         template = dataFile['template']
         th = template[0]
         tl = template[1]
         return th, tl
     def read_file(filename):
         dataFile = h5py.File(filename, 'r')
         dqInfo = dataFile['quality']['simple']
         qmask = dqInfo['DQmask'][...]
         meta = dataFile['meta']
         gpsStart = meta['GPSstart'][()]
         utc = meta['UTCstart'][()]
         duration = meta['Duration'][()]
         strain = dataFile['strain']['Strain'][()]
         dt = (1.0 * duration) / len(strain)
         dataFile.close()
         return strain, dt, utc
```

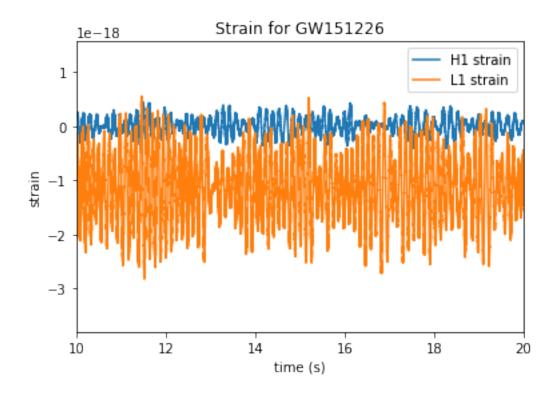
```
[3]: # Loading files so the data from the same events have the same index
with open(datadir + 'BBH_events_v3.json') as json_file:
    bbhev = json.load(json_file)
```

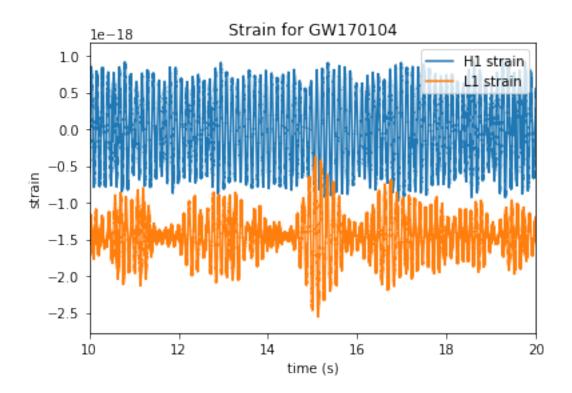
```
ths = []
tls = []
hstrain = []
hdt = []
hutc = []
lstrain = []
ldt = []
lutc = []
events = list(bbhev.keys())
for event in events:
    th, tl = read_template(datadir + bbhev[event]['fn_template'])
    ths.append(th)
    tls.append(tl)
    s, t, u = read_file(datadir + bbhev[event]['fn_H1'])
    hstrain.append(s)
    hdt.append(t)
    hutc.append(u)
    s, t, u = read_file(datadir + bbhev[event]['fn_L1'])
    lstrain.append(s)
    ldt.append(t)
    lutc.append(u)
```

```
for i in range(len(events)):
    plt.figure()
    plt.plot(hdt[i] * np.arange(len(hstrain[i])), hstrain[i], label='H1 strain')
    plt.plot(ldt[i] * np.arange(len(lstrain[i])), lstrain[i], label='L1 strain')
    plt.xlabel('time (s)')
    plt.legend(loc='upper right')
    plt.title(f"Strain for {events[i]}")
    plt.xlim((10, 20))
    plt.ylabel('strain')
    plt.savefig(f"strain_{events[i]}.png")
```

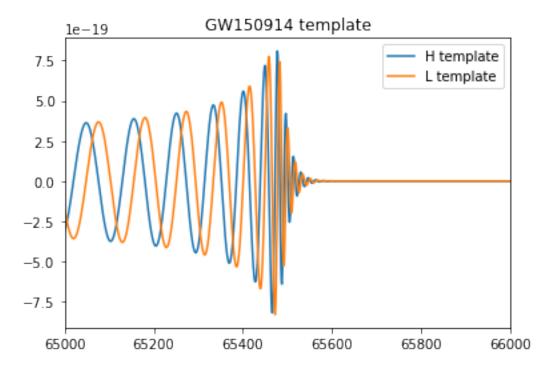


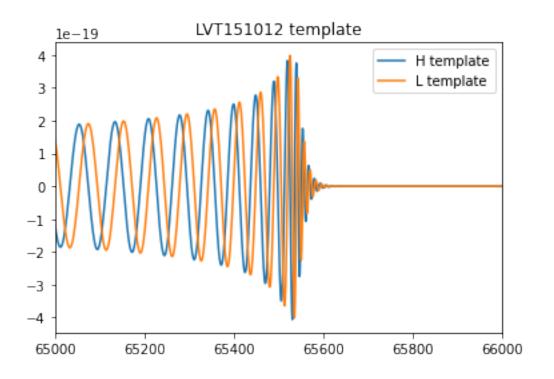


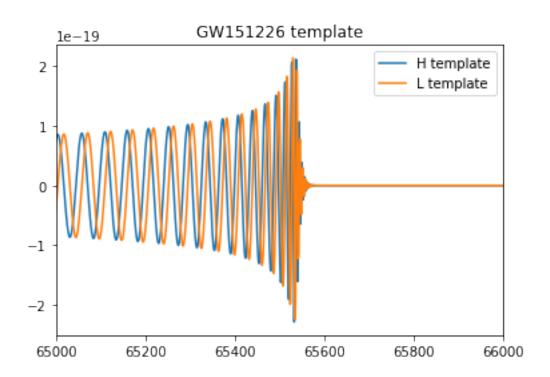


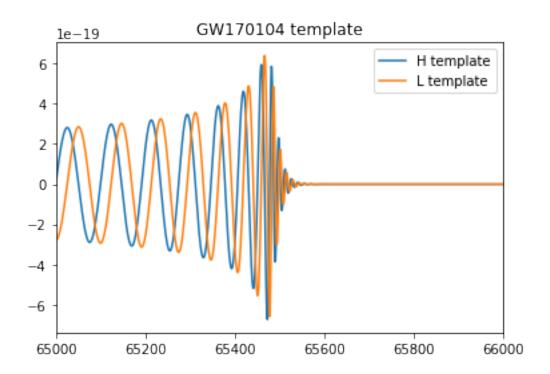


```
[5]: # Vizualizing templates
for i in range(len(events)):
    plt.figure()
    plt.plot(ths[i], label='H template')
    plt.plot(tls[i], label='L template')
    plt.legend(loc='upper right')
    plt.title(f"{events[i]} template")
    plt.xlim((6.5e4, 6.6e4))
    plt.show()
```



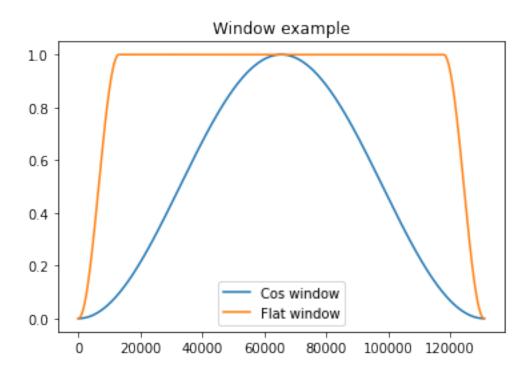






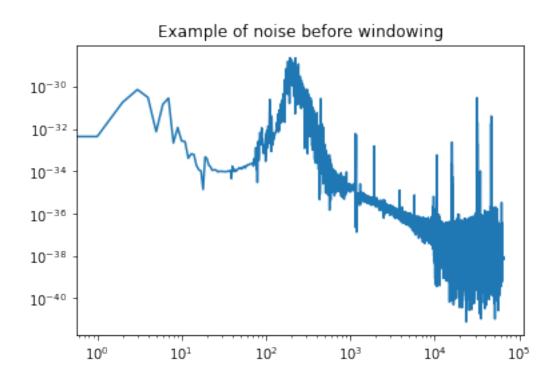
I used window function that was suggested in class (make\_flat\_window). It looks like simplified version of Tukey window, which is good for this case, because we are interested in obtaining wide noise range in the final result.

```
[6]: def make_window(n):
         x = np.linspace(-np.pi, np.pi, n)
         return 0.5 + 0.5 * np.cos(x)
     def make_flat_window(n, m):
         tmp = make_window(m)
         win = np.ones(n)
         mm = m//2
         win[:mm] = tmp[:mm]
         win[-mm:] = tmp[-mm:]
         return win
    n = len(hstrain[0])
     win = make_window(n)
     win_flat = make_flat_window(n,n//5)
     plt.plot(win, label='Cos window')
     plt.plot(win_flat, label='Flat window')
     plt.legend()
     plt.title("Window example")
     plt.savefig('window_example.png')
```

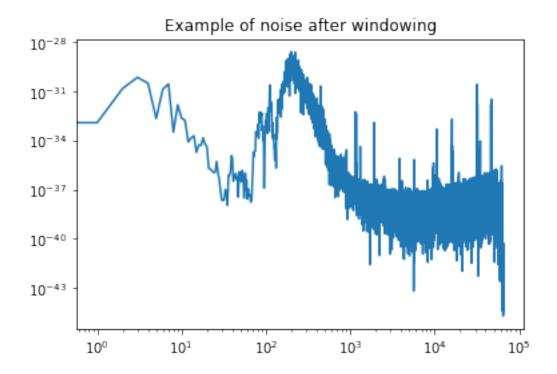


To make a noise model we treat all data as a noise and then smoothing it, so the actual signal we are interested in will remain a 'deviation' from the noise model. This 'deviation' is hidden somewhere in the small numerous peaks and smoothing the model we make sure that actual signal will not be removed during 'whitening' of data.

```
[7]: # Threat all data as a noise
sft = np.fft.rfft(hstrain[0])
Nft = np.abs(sft)**2
plt.loglog(Nft)
plt.title("Example of noise before windowing")
plt.savefig("noise_before_windowing.png")
```

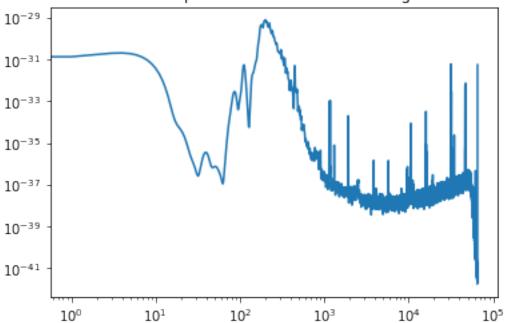


```
[8]: sft = np.fft.rfft(win_flat * hstrain[0])
Nft = np.abs(sft)**2
plt.loglog(Nft)
plt.title("Example of noise after windowing")
plt.savefig("noise_after_windowing.png")
```



```
[9]: # I tried using more sophisticated methods of smoothing including smoothing by \Box
     →convolving data with box function,
     # but in the end the result is almost the same in every case so I decided to
     →use the simpliest method suggested in class
     # def smooth_convolve(y, box_pts):
           box = np.ones(box_pts) / box_pts
           y_smooth = np.convolve(y, box, mode='same')
           return y_smooth
     def smooth(y):
         for i in range(10):
            y = (y + np.roll(y, 1) + np.roll(y, -1)) / 3
         return y
     Nft = smooth(Nft)
     plt.loglog(Nft)
     plt.title("Example of noise after windowing")
    plt.savefig("noise_after_smoothing.png")
```

## Example of noise after windowing



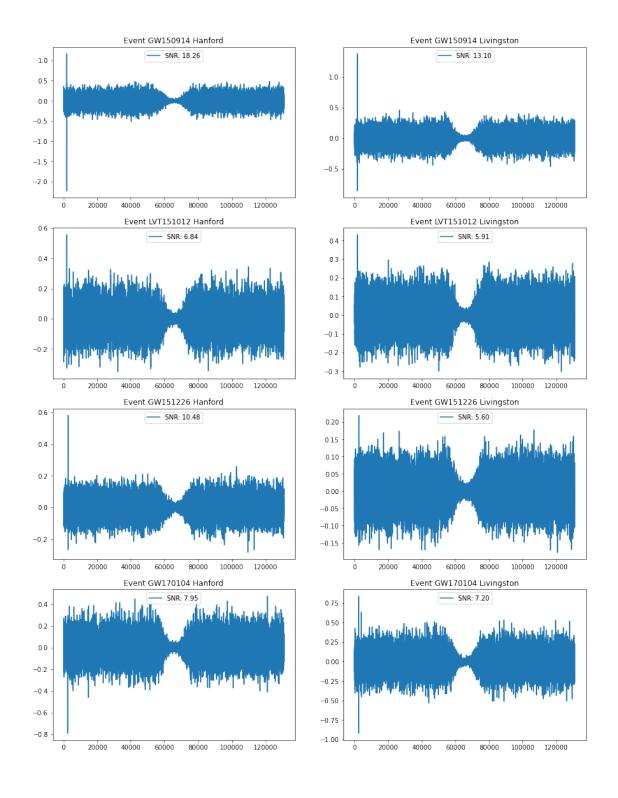
```
[10]: # Calculating noise model for each observation: windowing and smoothing
      hnm = [smooth(np.abs(np.fft.rfft(make_flat_window(len(s), len(s)//5) * s))**2)__
      →for s in hstrain]
      lnm = [smooth(np.abs(np.fft.rfft(make_flat_window(len(s), len(s)//5) * s))**2)__
      →for s in lstrain]
      # 'Whitening' data
      hsft_white = [np.fft.rfft(make_flat_window(len(s), len(s)//5) * hstrain[i]) /__
      →np.sqrt(hnm[i]) for i in range(len(hstrain))]
      lsft white = [np.fft.rfft(make flat window(len(s), len(s)//5) * lstrain[i]) /___
       →np.sqrt(lnm[i]) for i in range(len(lstrain))]
      # Windowing and 'whitening' templates
      thsft_white = [np.fft.rfft(ths[i] * make_flat_window(len(ths[i]), len(ths[i])//
      →5)) / np.sqrt(hnm[i]) for i in range(len(events))]
      tlsft_white = [np.fft.rfft(tls[i] * make_flat_window(len(tls[i]), len(tls[i])//
      →5)) / np.sqrt(lnm[i]) for i in range(len(events))]
      # Back to real space
      ths_white = [np.fft.irfft(tft) for tft in thsft_white]
      tls_white = [np.fft.irfft(tft) for tft in tlsft_white]
      # Matched filters!
```

```
fig, axs = plt.subplots(4, 2, figsize=(15, 20))

for i in range(len(events)):
    axs[i, 0].plot(hxcorr[i], label=f'SNR: {hsnr[i]:.2f}')
    axs[i, 0].title.set_text(f'Event {events[i]} Hanford')
    axs[i, 0].legend(loc='upper center')

for i in range(len(events)):
    axs[i, 1].plot(lxcorr[i], label=f'SNR: {lsnr[i]:.2f}')
    axs[i, 1].title.set_text(f'Event {events[i]} Livingston')
    axs[i, 1].legend(loc='upper center')

plt.savefig("matched_filters.png", dpi=500)
```

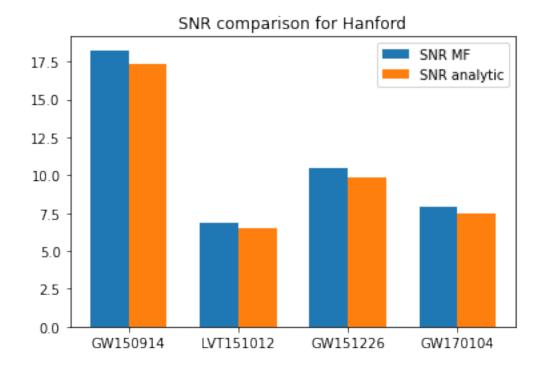


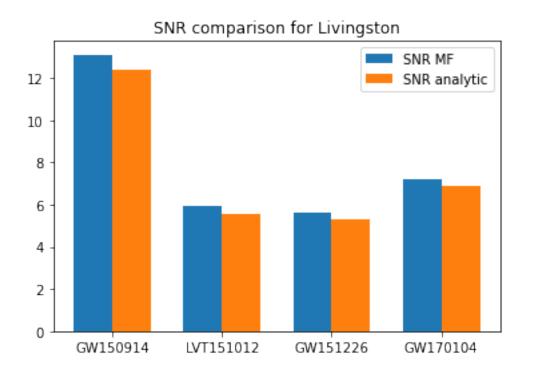
```
[13]: print("Combined SNR:")
for i in range(len(events)):
    print(f"{events[i]}:\t{np.sqrt(hsnr[i]**2 + lsnr[i]**2):.2f}")
```

Combined SNR:
GW150914: 22.47
LVT151012: 9.04
GW151226: 11.88
GW170104: 10.73

Comparison of measured and analytic SNR shows that thet are very close, but it is obvious that there is some bias because the analytic SNR is always lower than measured. It is caused, probably, by my method of calculation of analytic noise - I calculate it over whole model including the peak, which implies higher sigma and lower SNR.

```
[14]: # Analityc SNR estimation
      sig_ht = [np.std(tw) for tw in ths_white]
      sig_lt = [np.std(tw) for tw in tls_white]
      hsnr_teor = [np.max(np.abs(hxcorr[i])) / sig_ht[i] for i in range(len(events))]
      lsnr_teor = [np.max(np.abs(lxcorr[i])) / sig_lt[i] for i in range(len(events))]
      width = 0.35
      ind = np.arange(len(events))
      fig = plt.figure()
      ax = fig.add_subplot(111)
      bars1 = ax.bar(ind, hsnr, width, label='SNR MF')
      rects2 = ax.bar(ind + width, hsnr_teor, width, label='SNR analytic')
      ax.set_xticks(ind + width / 2)
      ax.set_xticklabels(events)
      plt.title("SNR comparison for Hanford")
      plt.legend()
      plt.savefig("cnr_comp_h.png")
      fig = plt.figure()
      ax = fig.add_subplot(111)
      bars1 = ax.bar(ind, lsnr, width, label='SNR MF')
      rects2 = ax.bar(ind + width, lsnr_teor, width, label='SNR analytic')
      ax.set_xticks(ind + width / 2)
      ax.set_xticklabels(events)
      plt.title("SNR comparison for Livingston")
      plt.legend()
      plt.savefig("cnr_comp_l.png")
```





```
[16]: # Finding half-frequency
     for i in range(len(events)):
         print(f"Event:\t{events[i]}")
         powh = np.abs(thsft_white[i])**2
         powl = np.abs(tlsft_white[i])**2
         normh = np.sum(powh)
         norml = np.sum(powl)
         cum weighth = np.cumsum(powh) / normh
         cum_weightl = np.cumsum(powl) / norml
         fridnh = np.searchsorted(cum weighth, 0.5)
         fridnl = np.searchsorted(cum_weightl, 0.5)
         freqsh = np.fft.rfftfreq(len(powh), d=hdt[i])
         freqsl = np.fft.rfftfreq(len(powl), d=ldt[i])
         print("\t|| Half cumsum index check:")
         print('\t||', cum_weighth[fridnh])
         print('\t||', cum_weightl[fridnl])
         print(f"H half-freq:\t{freqsh[fridnh]}")
         print(f"L half-freq:\t{freqsl[fridnl]}")
         print("======"")
     Event: GW150914
             || Half cumsum index check:
             11 0.5000247285568483
             | | 0.5000080829513244
     H half-freq:
                    202.99690251308422
     L half-freq:
                    229.9339914857256
     _____
     Event: LVT151012
             || Half cumsum index check:
             | | 0.5003575651317228
             | | 0.5000241093983616
     H half-freq:
                    152.93516639455575
     L half-freq:
                    193.37204937668798
     Event: GW151226
             | | Half cumsum index check:
             | | 0.5000928518939606
             || 0.5002357457707245
     H half-freq:
                    148.4352350580588
     L half-freq:
                    215.12171750308985
     _____
     Event: GW170104
             || Half cumsum index check:
             11 0.5000059654955388
             || 0.5000067909296776
```

H half-freq:

L half-freq:

188.9971161328715

150.24770740192562

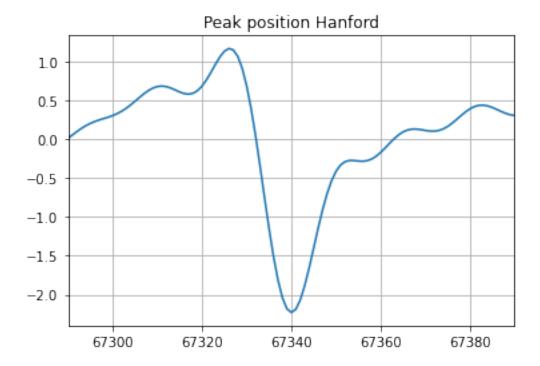
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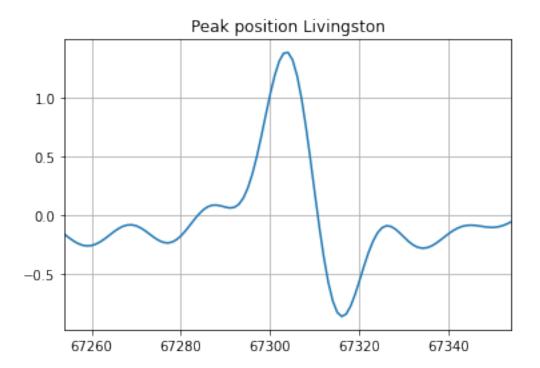
I will estimate localization error on the example of one event

```
[88]: h_peak_pos = np.argmax(np.abs(np.fft.fftshift(hxcorr[0])))
    plt.plot(np.fft.fftshift(hxcorr[0]))
    plt.xlim((h_peak_pos-50, h_peak_pos+50))
    plt.title("Peak position Hanford")
    plt.grid()
    plt.show()

l_peak_pos = np.argmax(np.abs(np.fft.fftshift(lxcorr[0])))
    plt.plot(np.fft.fftshift(lxcorr[0]))
    plt.xlim((l_peak_pos-50, l_peak_pos+50))
    plt.title("Peak position Livingston")
    plt.grid()
    plt.show()

print("Time delay:\t", (h_peak_pos - l_peak_pos) * hdt[0])
    print("Error of t is approximately:\t", 10 * hdt[0])
```





Time delay: 0.0087890625

Error of t is approximately: 0.00244140625

The FWHM of the peak was estimated just looking at the plot and equals approx. 10 time intervals or 0.00244140620 seconds. Time difference between two peaks is 0.0087890625. All further estimations are in the part\_f.pdf.