```
import numpy as np
              import matplotlib.pyplot as plt
             Problem 1
             Show that a Poisson distribution converges to a Gaussian in the large \lambda limit (use Stirling and first
             Poisson distribution: P(k,\lambda)=rac{e^{-\lambda}\lambda^k}{k!} , Gaussian distribution:
             G(x,\mu,\sigma)=rac{1}{\sqrt{2\pi\sigma}}exp(-rac{(x-\mu)^2}{2\sigma^2}) .
             Write P = P(x, \lambda) and take the log:
             ln(P(x)) = ln(e^{-\lambda}) + ln(\lambda^x) - ln(x!)
             Use Stirling's approximation: n! pprox e^{-n} n^n \sqrt{2\pi n} :
             ln(P(x)) = -\lambda + x ln(\lambda) - (ln(e^{-x}x^x\sqrt{2\pi x}))
             ln(P(x)) = -\lambda + x ln(\lambda) + x - x ln(x) - rac{1}{2} ln(2\pi x)
             ln(P(x)) = (x - \lambda) - x ln(x/\lambda) - \frac{1}{2} ln(2\pi x)
             Let y=x-\lambda , then \frac{x}{\lambda}=1+\frac{y}{\lambda} :
             ln(P(y)) = y - (y+\lambda)ln(1+rac{y}{\lambda}) - rac{1}{2}ln(2\pi(y+\lambda))
             For small \epsilon, the log expansion is ln(1+\epsilon)pprox \epsilon+rac{\epsilon^2}{2} . Since \lambda is large, rac{y}{\lambda} is very small. So
             expand ln(1+\frac{y}{\lambda}):
             ln(P(y)) = y - (y+\lambda)(rac{y}{\lambda} + rac{y^2}{2\lambda^2}) - rac{1}{2}ln(2\pi(y+\lambda))
             Eliminate the 1/\lambda^2 term because it will approach zero for large \lambda.
             ln(P(y)) = y - (y + \lambda)(rac{y}{\lambda}) - rac{1}{2}ln(2\pi(y + \lambda))
             ln(P(y)) = y - (rac{y^2}{\lambda} + y) - rac{1}{2}ln(2\pi(y+\lambda))
             ln(P(y)) = rac{y^2}{\lambda} - rac{1}{2}ln(2\pi(y+\lambda))
             Take the exponent:
             P(y) = rac{1}{\sqrt{2\pi\lambda}} exp(-rac{y^2}{2\lambda})
             Sub x=y, \lambda=\sigma^2:
             P(x) = \frac{1}{\sqrt{2\pi}} exp(\frac{x^2}{2\lambda}) = G(x) (Gaussian centered at zero).
             Problem 2
             5\sigma is the gold standard for a believable result. Define the Gaussian approximation as "good
             enough" if it agrees with the Poisson distribution to within a factor of 2. How large does n need to
             The premise is that P(x=\lambda+c\lambda)=2G(x=\lambda+c\lambda) , where c is a constant (3 or 5).
             Set the constants in the Gaussian in terms of \lambda: \mu = \lambda , \sigma^2 = \lambda :
             Then P(x,\lambda)=rac{e^{\lambda}\lambda^x}{x!} and 2G(x,\lambda)=rac{2}{\sqrt{2\pi\lambda}}exp(-rac{(x-\lambda)^2}{2\lambda})=\sqrt{rac{2}{\pi\lambda}}exp(-rac{(x-\lambda)^2}{2\lambda})
             Then we solve for x in \frac{e^{\lambda}\lambda^x}{x!}=\sqrt{\frac{2}{\pi\lambda}}exp(-\frac{(x-\lambda)^2}{2\lambda}).
             Take the log of both sides and use Stirling's approximation for the x!.
             Left hand side:
             ln(LHS) = -\lambda + xln(\lambda) - ln(x!)
             ln(LHS) = -\lambda + xln(\lambda) - (-x + xln(x) + rac{1}{2}ln(2\pi x))
             ln(LHS) = x - \lambda + x ln(rac{\lambda}{x}) - rac{1}{2} ln(2\pi x)
             Right hand side:
             ln(RHS) = \frac{1}{2}ln(\frac{2}{\pi\lambda}) - \frac{(x-\lambda)^2}{\sqrt{2\lambda}}.
             The x (or n) where ln(LHS) = ln(RHS) is where these two curves meet, evaluated at
                   (X_{i}, x_{i}) = (X_{i}, \frac{1}{2} x_{i}, x_{i}) + (x_{i}, \frac{1}{2} x_{i
In [2]: \# define functions for ln(LHS) and ln(RHS) from the above equations:
             def log_lhs(x,L):
                    # returns log(Poisson(x,lambda))
                    return x - L + x*np.log(L/x) - 0.5*np.log(2*x*np.pi)
             def log_rhs(x,L):
                    # returns log(2*Gaussian(x,lambda))
                    return 0.5*np.log(2/(L*np.pi)) - ((x-L)**2/(2*L))
             # evaluate by taking x-->lamda+3*sqrt(lambda)
             L = np.linspace(0.1, 10, 1001)
             X3 = L + 3*np.sqrt(L)
             log_lhs_3 = log_lhs(X3,L)
             log_rhs_3 = log_rhs(X3, L)
             # find where sign of the difference between the right and left hand
             # sides change to determine n. This is where Poisson and Gaussian cross
             start_sign = np.sign(log_lhs_3[0]-log_rhs_3[0])
             # condition for if max lambda (L) isn't big enough:
             if np.sign(log_lhs_3[-1]-log_rhs_3[-1]) == start_sign:
                    print('extend L for 3 sigma')
             for i in np.arange(1,len(X3)):
                    if np.sign(log_lhs_3[i]-log_rhs_3[i]) != start_sign:
                          print(r'Poisson = Gaussian when n = \{\} for 3 sigma'.format(int(X))
             3[i])))
                          break
             # plot the results
             plt.figure()
             plt.plot(X3, log_lhs_3, label='log(lhs)')
             plt.plot(X3, log_rhs_3, label='log(rhs)')
             plt.axvline(n3, c='k', label='n')
             plt.legend()
             plt.grid()
             # Repeat:
             # evaluate by taking x-->lambda+5*sqrt(lambda)
             L = np.linspace(0.1,600,1001)
             X5 = L + 5*np.sqrt(L)
             log_lhs_5 = log_lhs(X5, L)
             log_rhs_5 = log_rhs(X5,L)
             # find where sign of the difference between the right and left hand
             # sides change to determine n. This is where Poisson and Gaussian cross
             start\_sign = np.sign(log\_lhs\_5[0]-log\_rhs\_5[0])
             # condition for if max lambda (L) isn't big enough:
             if np.sign(log_lhs_5[-1]-log_rhs_5[-1]) == start_sign:
                    print('extend L for 5 sigma')
             for i in np.arange(1,len(X5)):
                    if np.sign(log_lhs_5[i]-log_rhs_5[i]) != start_sign:
                          n5 = X5[i]
                          print(r'Poisson = Gaussian when n = {} for 5 sigma'.format(int(X))
             5[i])))
                          break
             # plot results
             plt.figure()
             plt.plot(X5, log_lhs_5, label='log(lhs)')
             plt.plot(X5, log_rhs_5, label='log(rhs)')
             plt.axvline(n5, c='k', label='n')
              plt.legend()
             \beta q i s_{\beta} \rho \tilde{n}_{d7} \hat{s}áussian when n = 17 for 3 sigma
             Poisšon = Gaussian when n = 696 for 5 sigma
               -2.5
               -3.0
               -3.5
               -4.0
               -4.5
               -5.0
                          log(lhs)
               -5.5
                        log(rhs)
               -6.0
                                  5.0
                                          7.5
                                                 10.0
                                                         12.5
                                                                 15.0
                                                                         17.5
                                                  log(lhs)
                                                log(rhs)
                -8
               -10
               -12
               -14
               -16
                             100
                                     200
                                             300
                                                     400
                                                             500
                                                                      600
                                                                              700
             Fig: The Gaussian distribution is in agreement with the Poisson distribution within 5\sigma when
             n=696 . The Gaussian is in agreement with Poisson within 3\sigma when n=17 .
             Problem 3
             n Gaussian-distributed data points with identical \sigma and unknown mean. What is the error on the
             maximum-likelihood estimate of the mean? If half the data is wrong by a factor of \sqrt{2} (variance is
             off by a factor of 2), what is the true error on the estimated mean, compare it to maximum-
             likelihood with correct noise estimates. What if 1% of the data is underweighted by a factor of
             This derivation follows from Worked Example 2 in Lecture 2.
             For n Gaussian-distributed data points we have the model m, data d, weights w, we have
             m=rac{\sum w_i d_i}{\sum w_i d_i} and the variance is rac{1}{\sum w_i}. (Sums are from i=1 to N)
             We define the weights as w_i = rac{1}{\sigma_i^2} .
             The error (E) is the square root of the variance:
            E=\sqrt{rac{1}{\sum w_i}}=\sqrt{rac{1}{\sum rac{1}{\sigma_i^2}}}
             Since \sigma_i = \sigma (same standard deviation for all data points):
             \sum_i^N rac{1}{\sigma_i^2} = Nrac{1}{\sigma^2} .
             The error is then:
             E=\sqrt{rac{\sigma^2}{N}}=rac{\sigma}{\sqrt{N}} .
             If half of the data are wrong, then the error is:
             E = (\sum_i^{N/2} rac{1}{\sigma_i^2} + \sum_i^{N/2} rac{1}{2\sigma_i^2})^{-1/2} = (rac{N}{2\sigma^2} + rac{N}{2\sigma^2})^{-1/2} = (rac{6N}{8\sigma^2})^{-1/2} = rac{2\sigma}{\sqrt{3N}} pprox 1.15 rac{\sigma}{\sqrt{N}}
             This is about 15% off from the error on the maximum-likelihood estimated mean.
             If 1% of the data are underweighted by a factor of 100, this means w_i 	o w_i/100 or
             \sigma_i 
ightarrow 100\sigma_i . The error is then:
             E = (\sum_i^{N/100} rac{1}{100\sigma_i^2} + \sum_i^{99N/100} rac{1}{\sigma_i^2})^{-1/2} = (rac{N}{100^2\sigma^2} + rac{99N}{100\sigma^2})^{-1/2} = (rac{9901}{10000})^{-1/2} rac{\sigma}{\sqrt{N}} pprox 1.00
             This is about 0.5% off from the error on the maximum-likelihood estimated mean.
             If 1% of the data are overweighted by a factor of 100, this means w_i	o 100w_i or \sigma_i	o rac{\sigma_i}{100} .
             The error is then:
             E = (\sum_i^{N/100} rac{100}{\sigma_i^2} + \sum_i^{99N/100} rac{1}{\sigma_i^2})^{-1/2} = (rac{100N}{100\sigma^2} + rac{99N}{100\sigma^2})^{-1/2} = (rac{199}{100})^{-1/2} rac{\sigma}{\sqrt{N}} pprox 0.701 rac{\sigma}{\sqrt{N}}
             This is about 30% off from the error on the maximum-likelihood estimated mean.
In [3]: # do the number math:
             2/np.sqrt(3), (9901/10000)**-0.5, (199/100)**-0.5
Out[3]: (1.1547005383792517, 1.004987059618685, 0.708881205008336)
             Problem 4
             Write a program that generates Gaussian noise + a signal (unit Gaussian). Estimate noise by
             assuming it's constant within a chunk and equal to the scatter in the observed data. Show that the
             least-squares estimate for one chunk is unbiased, but least-squares estimate for many chunks is
             biased LOW. Fit an amplitude and error to each chunk and use that to get an overall amplitude
In [4]: # set number of points, size of individual chunks, size of multiple chun
             npts, chunk_size, chunk_size_new = 51,3,17
             # this was for a test with more points:
             #npts, chunk_size, chunk_size_new = 10000,10,500
             # define x-values
             x = np.linspace(-5, 5, npts)
             # generate the signal+noise
             data = np.exp(-x**2/2) + np.random.randn(len(x))
              # define size of chunks
             nchunks = int(len(x)/chunk_size)
             # print a sanity check:
              #print('individual chunks have size = {}, for a total of {} chunks. Tota
             1 points is {}={}'.format(chunk_size,nchunks,npts, chunk_size*nchunks))
             # divide x into chunks
             x_chunks = [x[chunk_size*i:chunk_size*(i+1)] for i in range(nchunks)]
             data_chunks = [data[chunk_size*i:chunk_size*(i+1)] for i in range(nchunk
             s)]
             # estimate constant noise for each chunk
             noise_chunk = [np.std(data_chunks[i]) for i in range(nchunks)]
             # repeat so that each data point has an associate noise
             noise = np.repeat(noise_chunk,chunk_size)
             # least-squares estimate for each chunk (should be unbiased):
             amp_fits, err_fits = np.zeros(nchunks), np.zeros(nchunks)
             for i in range(nchunks):
                    # take one chunk of data:
                    patch = data_chunks[i]
                    # model is the template signal:
                    mymodel = np.exp(-patch**2/2)
                    # the least-squares best-fit amplitude for constant noise:
                    amp_fit = np.sum(mymodel*patch)/np.sum(mymodel**2)
                    amp_fits[i] = amp_fit
                    # calculate the error (scatter between data and the model):
                    err_fit = np.std(patch - mymodel*amp_fit)
                    err_fits[i] = err_fit
             # least-squares estimate for multiple chunks (should be biased low):
             nchunks_new = int(len(x)/chunk_size_new) # look at more chunks
             # print a sanity check:
             #print('multiple chunks have size = {}, for a total of {} chunks. Total
              points is {}={}'.format(chunk_size_new,nchunks_new,npts, chunk_size_new
              *nchunks_new))
             amp_fits_multip, err_fits_multip = np.zeros(nchunks_new), np.zeros(nchun
             ks_new)
             for i in range(nchunks_new):
                    patch = data[chunk_size_new*i:chunk_size_new*(i+1)]
                    # model is the template signal:
                    mymodel = np.exp(-patch**2/2)
                    # get the least-squares best-fit amplitude for constant noise:
                    amp_fit = np.sum(mymodel*patch)/np.sum(mymodel**2)
                    amp_fits_multip[i] = amp_fit
                    # calculate the error (scatter between data and the model):
                    err_fit = np.std(patch - mymodel*amp_fit)
                    err_fits_multip[i] = err_fit
             # use weighted average to get overall amplitude and error:
             amp_avg = np.sum(amp_fits)/len(amp_fits)
             err_avg = np.sum(err_fits)/len(err_fits)
             # plot the results
             plt.figure()
             plt.scatter(amp_fits, err_fits, marker='.', label='individual chunks')
             plt.scatter(amp_fits_multip, err_fits_multip, marker='.', label='multipl
              e chunks'
Out[4]: pmatelattab(lmgendgLegendaat, 0mafceadge859abel='average')
             plt.ylabel('fitted error')
             plt.x
                                                                   individual chunks
             p1t<sup>1.6</sup>1
                                                                   multiple chunks
                                                                   average
                 1.4
                 1.2
              10 0.8
```

jointly analyzing multiple seem to have higher fitted errors (meaning this analysis is biased high? I **Problem 5** 

Fig: The fitted amplitudes are unbiased for the individual chunks, while the fitted amplitudes from

0.75 1.00

0.8

0.6 0.4

Show that in rotation space,  $N_{ij} = \langle n_i n_j \rangle$ . Original expression for  $\chi^2$  is  $\chi^2=n^TN^{-1}n$  , where n are the residuals between the model and

-0.75 -0.50 -0.25 0.00

0.25

the data: n=a-A(m) , and N is diagonal matrix with  $N_{ii}=< n_i^2>$  .

fitted amplitude

0.50

```
Introduce orthogonal rotation matrix V where VV^T=V^TV=I , which does not change \chi^2:
\chi^2 = n^t V^T V N^{-1} V^T V n = (V n)^T (V N V^T)^{-1} (V n) .
```

Let  $Vr = ilde{n}$  (so  $(Vr)^T = ilde{n}$ ),  $VNV^T = ilde{N}$  :

 $\chi^2 = ilde{n} ilde{N}^{-1}$   $\tilde{n}$  , where now  $ilde{N}$  is not diagonal. Use  $ilde{n}_i = V_{::i}^T n$  and  $ilde{n}_j = n^T V_{::j}$  to find  $< ilde{n_i} ilde{n_j}>$  :

 $<\tilde{n_i}\tilde{n_j}>=< V_{:,i}^T n n^T V_{:,j}>$  . Since  $nn^T=n^2$  and < nnT>=N :  $< V_{:,i}^T nn^T V_{:,j}>=V_{:,i}^T N V_{:,j}= ilde{N_{ij}}$  .

Therefore  $< ilde{n_i} ilde{n_j}> = ilde{N_{ij}}$  . In [ ]: