PHYS-512: Problem Set 5

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- 1. With the initial set of parameters, χ^2 results in roughly chisq = 15268. For 2501 degrees of freedom, I would expect a good fit model to result in $\chi^2 = 2501 \pm 71$ (or in general χ^2 should be at $n \pm \sqrt{2n}$). Clearly, χ^2 lies outside these bounds. With the new set of parameters, χ^2 is calculated to be chisq = 3272.2053559202122. I would again conclude that the fit is not acceptable, although a lot better.
- 2. To implement Newton's method for this problem, I left the numeric differentiator from the previous problem set practically unchanged. The key difference is that I implemented a stopping condition for this problem. The stopping condition is triggered whenever the χ^2 value of two successive trial steps is closer then a tolerance factor tol that I've fixed to be 0.5. Otherwise, Newton's method is executed for a number of steps iterator. As initial condition chi (the list that will store all later computed values for χ^2) is set to hold 4000.

```
[TRUNCATED]
iterator = 101
[TRUNCATED]
# starting condition
chi = [4000]
d_{chi} = []
tol = 0.5
converged = False
Ninv = np.eye(len(spec))/(errs**2)
for i in range(iterator):
   pred, grad = gradient(model_params[i])
    r = spec - pred
    r = np.matrix(r).transpose()
    grad = np.matrix(grad)
    lhs = grad.T*Ninv*grad
    rhs = grad.T*Ninv*r
    u, s, vh = np.linalg.svd(lhs)
    dm = vh.T@np.linalg.inv(np.diag(s))@u.T@rhs
    dm = dm.reshape(6.)
    if i == iterator - 1:
        break
```

new_chi = get_chs(pred, spec) chi.append(new_chi) converged = np.abs(chi[-2] - chi[-1]) < tol # print(f"new chi = {new_chi}") if converged: print('Newton Method has converged.') break print(f"Params in current loop: {model_params[i]}\nWith chi sq: {new_chi}.")</pre>

Doing this results in a χ^2 value of 2576.8160468697643 and parameters (dropped dimensions)

model_params[i + 1] = model_params[i] + dm

$$H_0 = 67.7$$
 $\Omega_b h^2 = 2.23 \times 10^{-2}$
 $\Omega_c h^2 = 0.1191$ $\tau = 7.0 \times 10^{-2}$
 $A_s = 2.16 \times 10^{-9}$ $n_s = 9.69$

with parameter errors

else:

$$\Delta H_0 = 0.5$$
 $\Delta \Omega_b h^2 = 0.02 \times 10^{-2}$ $\Delta \Omega_c h^2 = 0.0008$ $\Delta \tau = 3.2 \times 10^{-2}$ $\Delta h_s = 0.13 \times 10^{-9}$ $\Delta h_s = 0.006$

According to the reasoning in question 1, I would say that this is indeed an acceptable fit.

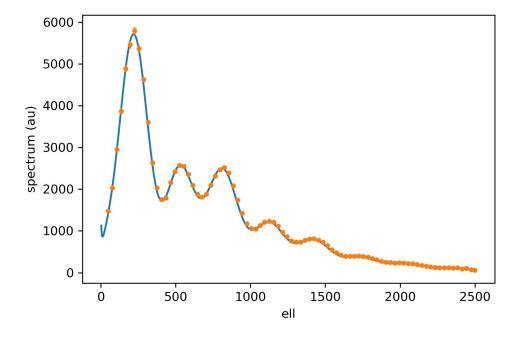


Figure 1: Fit to spectrum using Newton's method.

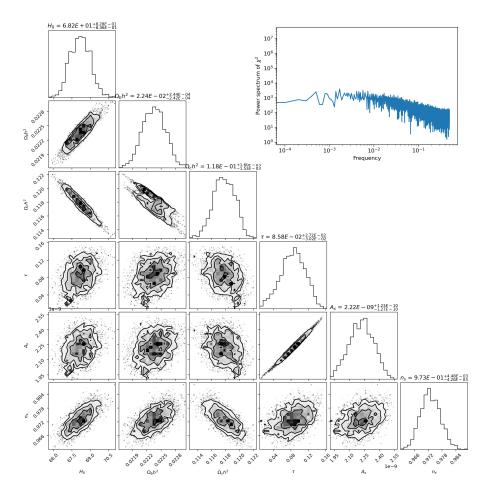


Figure 2: Corner plots for the first run of the MCMC chain with 10,000 steps. The inset shows a double-log plot of the power spectrum for χ^2 .

3. With the curvature matrix from the previous problem, I initialized parameters and step sizes in parameter space for the MCMC chain - They are imported with the call to newton.py as

```
from newton import curv, best, errs, spec, get_spectrum
```

The MCMC chain is set up to run for n_steps number of steps (in this case 10,000). If a step in the chain is accepted, the new parameters and the associated value for χ^2 is saved in separate textfiles by the call

```
np.savetxt('mcmc_params.txt', par)
np.savetxt('chi_square.txt', chi_sq)
```

The results are shown in Fig. 2. One can see that the parameter values are more or less normally distributed. The FFT power spectrum for χ^2 also shows this - The frequency components are more or less constant with a linear decrease at higher frequencies. With $\Omega_b = 4.82 \pm 0.17$ and $\Omega_c = 25.7 \pm 0.96$, I get $\Omega_{\Lambda} = -29.5 \pm 1.1$

Complete Code:

```
import numpy as np
from newton import curv, best, errs, spec, get_spectrum
def get_chi_sq(data, params):
    data = get_spectrum(params)[:len(spec)]
    chisq = np.sum((data - spec)**2/errs**2)
    return chisq
n_steps = 10000
steps_taken = 0
chi_sq = []
par = []
# initialize parameter list by using best fit +/- a little something
init_par = np.random.multivariate_normal(best, np.linalg.inv(curv)/2)
chi_sq.append(get_chi_sq(spec, init_par))
par.append(init_par)
while steps_taken < n_steps:</pre>
    # generate new parameters
    step = np.random.multivariate_normal(np.zeros(best.size),
                np.linalg.inv(curv)/2)
    par_new = par[-1] + step
    # calculate new chi squared
   new_chi = get_chi_sq(spec, par_new)
    delta_chi = new_chi - chi_sq[-1]
    prob_step = np.exp(-0.5*delta_chi)
    # if prob_step is greater then one, chi will have increased
    # and we want to discard that step, otherwise accept with a certain
    # probability:
    if prob_step > np.random.rand(1):
        par.append(par_new)
        chi_sq.append(new_chi)
    else:
        # if step is not taken, copy last entries for params and chi_sq
        par.append(par[-1])
        chi_sq.append(chi_sq[-1])
    steps_taken += 1
    np.savetxt('mcmc_params.txt', par)
   np.savetxt('chi_square.txt', chi_sq)
    if steps_taken % 100 == 0:
        print(f"At {steps_taken/n_steps*100:.2f} percent")
```

4. I first sampled the latter half of my prior MCMC run to get an importance sampled estimate for the new parameters and the covariance matrix. I calculated the weights according to the rule

$$w = \exp\left(-0.5\delta\chi^2\right) = \exp\left(-0.5\left(\frac{\tau - \tau_{\text{pol}}}{\sigma_{\text{pol}}}\right)^2\right)$$
 (1)

where the subscript pol identifies the polarization data values. The normalized weights are then used to generate a new set of parameters, sampled_params by averaging the last 5000 steps. Similarly for the covariance matrix.

```
import numpy as np
```

```
mcmc_params = np.loadtxt('mcmc_params.txt')
tau_pol = 0.054
tau unc = 0.0074
# first step: importance sampling
# selecting only last 5000 steps
mcmc_params = mcmc_params[5001:,:]
# get tau values
tau = mcmc_params[:,3]
# define delta chi squared
delta_chi = (tau - tau_pol)**2/tau_unc**2
# define weights
w = np.exp(-0.5*delta_chi)
# normalize weights
w = w/np.sum(w)
# importance sample parameters
sampled_params = np.zeros(6)
for j in range(6):
    sampled_params[j] = sum(mcmc_params[:,j]*w)
# get covariance matrix
cov_mat = np.cov(mcmc_params, rowvar=False, aweights=w)
```

I get the following values for the parameters:

$$H_0 = 68.2$$
 $\Omega_b h^2 = 2.22 \times 10^{-2}$ $\Omega_c h^2 = 0.176$ $\tau = 5.6 \times 10^{-2}$ $n_s = 9.73$

with parameter errors

$$\Delta H_0 = 0.7$$
 $\Delta \Omega_b h^2 = 0.02 \times 10^{-2}$ $\Delta \Omega_c h^2 = 0.001$ $\Delta \tau = 0.7 \times 10^{-2}$ $\Delta A_s = 0.03 \times 10^{-9}$ $\Delta n_s = 0.004$

It's interesting to note that τ does not automatically assume the tau_pol value of 0.054, but instead to 0.056. My guess would be that this is caused by the fact that the chain seems to actually prefer a much larger value for τ of 0.086 (see corner plots Fig. 2), skewing the distribution to larger values.

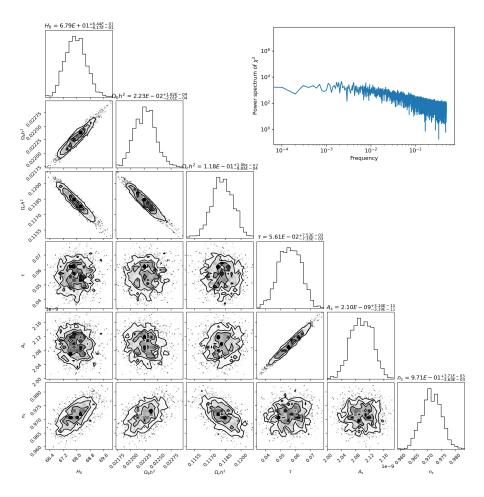


Figure 3: Corner plots for the second run of the MCMC chain with 10,000 steps. The inset shows a double-log plot of the power spectrum for χ^2 .

I modified my chi_sq getter function to include the constraint from the polarization data like so:

and run the chain again with the importance sampled parameters and covariances. Unfortunately, there seemed to be something wrong with my calculated curvature matrix. I instead used the same curvature matrix as in the previous problem, which should ideally not be too far off. The results are shown in Fig. 3. The distributions are again approximately normal, but I would say worse than in Fig. 2 and the chain is maybe not completely converged. Except for $\Omega_c h^2$, the parameters are approximately the same as the importance sampled ones, which is a little surprising considering that the importance sampled value should inherently account for the covariance between the two parameters. In both cases, I obtain a χ^2 value of around $\chi^2 = 2582.44$.