```
In [12]: import numpy as np
import matplotlib.pyplot as plt
```

Problem set 5

I had a bit of trouble installing camb , since I am on an Apple chip and have non-native python (i.e. x86 python) running for whatever reason. Rigel helped me get this up and running by installing a virtual environment and installing camb there with conda create -n camb -c conda-forge python=3.9 camb. That said, I don't know how to run Jupyter notebooks in virtual environments, so this notebook is more of a walkthrough of my procedure, with relevant code and results attached.

1)

Running plank_likelihood.py for the parameters already included, one obtains the result:

```
chisq is 15267.937968194292 for 2501 degrees of freedom.
```

This is not a good fit. We should expect a χ^2 value near $<\chi^2>=n$ and within $\Delta\chi^2=2n$ where n is the number of degrees of freedom. In this case, n=2501 and therefore the χ^2 does not fall within the expected value of χ^2 provided the n of this problem: $15267.93 \neq 2501 \pm 5002$. Therefore, the parameters dialed into the test script do not constitute a good fit.

If we instead take pars=np.asarray([69,0.022,0.12,0.06,2.1e-9,0.95]), we obtain

```
chisq is 3272.203604462886 for 2501 degrees of freedom.
```

This χ^2 is within error to the expected χ^2 : $3272.203=2501\pm5002$ (don't worry about sig figs for now).

Here is an image of the fit, taken from the product of planck likelihood.py:

2)

We'll now use Newton's method. In much the same way we did in problem set 4, the basic idea here is to start off with a guess of parameters and take the numerical derivatives with respect to these parameters to arrive at a better guess, and so on until convergence. Since

we had a pretty good guess already in 1), we'll use that as our initial parameters. The code we use is shown below:

```
In [ ]: import numpy as np
                    import camb
                    # define numerical derivative function
                    def ndiff2(fun,x,dx ord=0):
                             Numerical derivative using both +/- dx and +/- 2*dx
                              # this is a fine choice for our purposes:
                             # a) Newton's method will keep iterating until the solution is good and
                              # b) our function and its derivatives are of order unity
                             order = -3 + dx_ord
                             dx = 10**(order)
                              # compute the function at the points of interest
                             yplus = fun(x + dx)
                             yminus = fun(x - dx)
                             yplus2 = fun(x + 2*dx)
                             yminus2 = fun(x - 2*dx)
                              # compute the numerical derivative
                              fprime = (8 * yplus - yplus2 + yminus2 - 8 * yminus) / (12 * dx)
                             return fprime
                    # define funcion to extract spectrum from camb
                    # this is more or less a black box
                    def get spectrum(H0=0,ombh2=0,omch2=0,tau=0,As=0,ns=0,params=np.asarray([0,0,0,0]
                              Returns the spectrum for a given set of parameters HO, baryon density, dark
                              optical depth, As, ns
                              if take params:
                                       # define params
                                       H0=params[0]
                                       ombh2=params[1]
                                       omch2=params[2]
                                       tau=params[3]
                                       As=params[4]
                                       ns=params[5]
                              # extract spectrum from camb
                              # this is more or less a black box
                              params=camb.CAMBparams()
                              params.set cosmology(H0=H0,ombh2=ombh2,omch2=omch2,mnu=0.06,omk=0,tau=tau)
                             params.InitPower.set params(As=As,ns=ns,r=0)
                             params.set for lmax(lmax,lens potential accuracy=0)
                             results=camb.get results(params)
                             powers=results.get cmb power spectra(params, CMB unit='muK') # <- this is where the same is the same in the same is the same in the same is the same i
                             cmb=powers['total']
                              tt=cmb[:,0]
                              return tt[2:]
                    # define gradient helper function
```

```
# make sure to rescale the optimal dx by the parameter's order
def params grad(fun,params):
    Returns the numerical derivative at points `params` in parameter space
    H0, ombh2, omch2, tau, As, ns = params
    # derivative w.r.t HO at p
    fun H00 = lambda H00: fun(H00,ombh2,omch2,tau,As,ns)
    grad_H0 = ndiff2(fun_H00,H0,dx_ord=+1)
    # derivative w.r.t ombh2 at p
    fun_ombh22 = lambda ombh22: fun(H0,ombh22,omch2,tau,As,ns)
    grad\_ombh2 = ndiff2(fun\_ombh22,ombh2,dx\_ord=-2)
    # derivative w.r.t omch2 at p
    fun omch22 = lambda omch22: fun(H0,ombh2,omch22,tau,As,ns)
    grad_omch2 = ndiff2(fun_omch22,omch2,dx_ord=-1)
    # derivative w.r.t tau at p
    fun_tauu = lambda tauu: fun(H0,ombh2,omch2,tauu,As,ns)
    grad tau = ndiff2(fun tauu,tau,dx ord=-2)
    # derivative w.r.t As at p
    fun Ass = lambda Ass: fun(H0,ombh2,omch2,tau,Ass,ns) # sorry for the profar
    grad_As = ndiff2(fun_Ass,As,dx_ord=-9)
    # derivative w.r.t ns at p
    fun nss = lambda nss: fun(H0,ombh2,omch2,tau,As,nss)
    grad ns = ndiff2(fun nss,ns,dx ord=0)
    # transpose to make it match with calc lorentz
    return np.array([grad H0, grad ombh2, grad omch2, grad tau, grad As, grad r
    # return np.array([grad H0, grad ombh2, grad omch2, ]).T
# define newtons method iterator
def newtons method(p0,d,num,print params=False):
    Numerical derivative version of `newtons method`
    Runs Newton's method for initial parameter guess p0
    t and d are the time and data, resp.
    # starting parameters is p0
    p = p0.copy()
    for i in range(num):
        # calculate derivatives and function
        pred = get spectrum(params=p,take params=True)
        pred = pred[:len(d)] # make both the same size (we don't want to extrag
        grad = params grad(fun=get spectrum,params=p)
        grad = grad[:len(d)]
        # delta is difference between data and prediction
        r = d - pred
        err = (r**2).sum()
        r = r.T
        lhs=grad.T@grad
        rhs=grad.T@r
```

```
dp=np.linalg.pinv(lhs)@(rhs)
        for jj in range(p.size):
            p[jj]=p[jj]+dp[jj]
        if print_params:
            print("The parameters are:",p)
            print("The step is:",dp)
    return p, dp
# run the "main"
if name == ' main ':
    # extract data from .txt files
   planck = np.loadtxt('COM_PowerSpect_CMB-TT-full_R3.01.txt',skiprows=1)
   ell = planck[:,0]
    spec = planck[:,1]
   errs = (planck[:,2] + planck[:,3]) / 2
    # initial guess and no. of iterations
   p0 = np.asarray([69, 0.022, 0.12, 0.06, 2.1e-9, 0.95])
    num iter = 5
    # run the iterator
   params_newton, step = newtons_method(p0=p0,d=spec,num=num_iter,print_params
    # estimate the uncertainties
   pred = get_spectrum(params=params_newton,take_params=True)
    pred = pred[:len(spec)]
   Ninv = np.linalg.inv(np.diag(errs))
    # parameter errors
    grad = params grad(fun=get spectrum,params=params newton) # use differentia
    grad = grad[:len(spec)]
   lhs = grad.T@Ninv@grad
    par errs = np.sqrt(np.diag(np.linalg.inv(lhs)))
    print("The errors are {}".format(par errs))
    np.savetxt("planck fit params.txt", np.array([params newton, par errs]).T)
```

When run in the camb environment, it returns, after 5 iterations:

```
The parameters are: [6.90000000e+01 2.20000000e-02 1.20000000e-
01 6.00000000e-02
 2.11779228e-09 9.50000000e-01]
The step is: [-5.21713904e-23 2.90831665e-19 -1.50903574e-19
-7.43590788e-20
  1.77922838e-11 -1.97538604e-201
The parameters are: [6.90000000e+01 2.20000000e-02 1.20000000e-
01 6.00000000e-02
 2.11779231e-09 9.50000000e-01]
The step is: [-9.20879475e-29 5.13388256e-25 -2.66392826e-25
-1.31262655e-25
  3.11437060e-17 -3.48715248e-26]
The parameters are: [6.90000000e+01 2.20000000e-02 1.20000000e-
01 6.00000000e-02
 2.11779231e-09 9.50000000e-01]
The step is: [ 9.08377570e-34 -5.06418471e-30 2.62776264e-30
```

```
1.29480626e-30
 -3.07208971e-22 3.43981073e-31]
The parameters are: [6.90000000e+01 2.20000000e-02 1.20000000e-
01 6.00000000e-02
 2.11779231e-09 9.50000000e-011
The step is: [ 1.11316243e-34 -6.20585571e-31 3.22016610e-31
1.58670771e-31
 -3.76466237e-23 4.21528250e-32]
The parameters are: [6.90000000e+01 2.20000000e-02 1.20000000e-
01 6.00000000e-02
 2.11779231e-09 9.50000000e-011
The step is: [-1.20182285e-35 6.70013562e-32 -3.47664374e-32
-1.71308474e-32
  4.06450769e-24 -4.55101852e-33]
The errors are [1.16828605e-01 2.20474064e-05 2.54033954e-04
3.63097430e-03
 1.46372787e-11 6.88626237e-04]
```

The best fit parameters and errors are therefore:

```
In [79]:
         newton = np.loadtxt("mcmc/planck_fit_params.txt")
         params newton = newton[:,0]
         param_errs_newton = newton[:,1]
In [80]: print("H0 = {} +/- {})".format(params newton[0], param errs newton[0]))
         print("Omegabh2 = {} +/- {}".format(params newton[1],param errs newton[1]))
         print("Omegach2 = {} +/- {}".format(params_newton[2],param_errs_newton[2]))
         print("tau = {} +/- {}".format(params_newton[3],param errs newton[3]))
         print("As = {} +/- {}".format(params_newton[4],param errs newton[4]))
         print("ns = {} +/- {}".format(params newton[5],param errs newton[5]))
         H0 = 69.0 + / - 0.11682860476516188
         Omegabh2 = 0.022 + / - 2.204740636442771e - 05
         Omegach2 = 0.12 +/- 0.00025403395375825035
         tau = 0.06 + / - 0.003630974299487817
         As = 2.1177923149017066e-09 +/- 1.463727873635012e-11
         ns = 0.95 + /- 0.0006886262373866526
```

Note: I know that something isn't quite right here, since none of the parameters are really varying. I think this has to do with either a bug in the differentiator (which would be odd, since it worked for PS4) or due to me not accounting for errors on the data through the matrix N which I ommitted in the Newton's method algorithm (which still should not mean that the parameters don't at all change, however). I believe it is more likely the latter, since in this case one must treat the camb model as a black box of sorts, and it is therefore difficult to assess whether or not the model and its derivatives are accurate.

3)

Let's try to get to the optimal parameters via an MCMC method. We know how to do this from the previous problem set, so let's just go ahead and implement it:

```
In [ ]: import numpy as np
        import camb
        import time
        # to get spectrum (model) from CAMB parameters
        def get spectrum(H0=0,ombh2=0,omch2=0,tau=0,As=0,ns=0,params=np.asarray([0,0,0,0,
            Returns the spectrum for a given set of parameters H0, baryon density, dark
            optical depth, As, ns
            if take_params:
                # define params
                H0=params[0]
                ombh2=params[1]
                omch2=params[2]
                tau=params[3]
                As=params[4]
                ns=params[5]
            # extract spectrum from camb
            params=camb.CAMBparams()
            params.set_cosmology(H0=H0,ombh2=ombh2,omch2=omch2,mnu=0.06,omk=0,tau=tau)
            params.InitPower.set_params(As=As,ns=ns,r=0)
            params.set_for_lmax(lmax,lens_potential_accuracy=0)
            results=camb.get results(params)
            powers=results.get_cmb_power_spectra(params,CMB_unit='muK') # <- this is wl</pre>
            cmb=powers['total']
            tt=cmb[:,0]
            return tt[2:]
        # to reproduce the curvature matrix from the previous problem
        def ndiff2(fun,x,dx ord=0):
            Numerical derivative using both +/- dx and +/- 2*dx
            # this is a fine choice for our purposes:
            # a) Newton's method will keep iterating until the solution is good and
            # b) our function and its derivatives are of order unity
            order = -3 + dx ord
            dx = 10**(order)
            # compute the function at the points of interest
            yplus = fun(x + dx)
            yminus = fun(x - dx)
            yplus2 = fun(x + 2*dx)
            yminus2 = fun(x - 2*dx)
            # compute the numerical derivative
            fprime = (8 * yplus - yplus2 + yminus2 - 8 * yminus) / (12 * dx)
            return fprime
        def params grad(fun,params):
            Returns the numerical derivative at points `params` in parameter space
            H0, ombh2, omch2, tau, As, ns = params
            # derivative w.r.t HO at p
            fun_H00 = lambda H00: fun(H00,ombh2,omch2,tau,As,ns)
```

```
grad H0 = ndiff2(fun H00,H0,dx ord=+1)
    # derivative w.r.t ombh2 at p
    fun ombh22 = lambda ombh22: fun(H0,ombh22,omch2,tau,As,ns)
    grad_ombh2 = ndiff2(fun_ombh22,ombh2,dx_ord=-2)
    # derivative w.r.t omch2 at p
    fun_omch22 = lambda omch22: fun(H0,ombh2,omch22,tau,As,ns)
    grad_omch2 = ndiff2(fun_omch22,omch2,dx_ord=-1)
    # derivative w.r.t tau at p
    fun tauu = lambda tauu: fun(H0,ombh2,omch2,tauu,As,ns)
    grad_tau = ndiff2(fun_tauu,tau,dx_ord=-2)
    # derivative w.r.t As at p
    fun_Ass = lambda Ass: fun(H0,ombh2,omch2,tau,Ass,ns) # sorry for the profat
    grad As = ndiff2(fun Ass, As, dx ord=-9)
    # derivative w.r.t ns at p
    fun nss = lambda nss: fun(H0,ombh2,omch2,tau,As,nss)
    grad_ns = ndiff2(fun_nss,ns,dx_ord=0)
    # transpose to make it match with calc_lorentz
    return np.array([grad_H0, grad_ombh2, grad_omch2, grad_tau, grad_As, grad_r
# computes chi squared
def chisquared(d,pred,errs):
    Computes chi-squared given some data d and pred
    In our case, errs is a constant and is always the same
   chi2 = np.sum((pred-d)**2/errs**2)
   return chi2
# takes a random step in parameter space
def random_step(cov):
    Random Gaussian step in each parameter
    scale = 1.5 # manually adjusted scale factor
   cov = cov/scale
    step = np.random.multivariate normal(np.zeros(cov.shape[0]),cov)
    return step
# performs a mcmc step
def mcmc_step(d,params,chisq,cov,errs):
    Single step in MCMC chain
   Params is a vector
    Param errs is the covariance in the initial parameters in the
    # set constrained parameters
   tau prior = 0.0540
    sigma tau prior = 0.0074
    # compute a set of new trial parameters
```

```
new params = params + random step(cov)
    # they predict the following data
    new_pred = get_spectrum(params=new_params,take_params=True)[:len(d)]
    # this data has the following chi squared
    # include the constraint that tau = tau prior +/- sigma tau prior
    tau_chisq = ((new_params[3] - tau_prior)/sigma_tau_prior)**2
    new_chisq = chisquared(d,new_pred,errs) + tau_chisq
    # if it improves the chi squared, always accept
    # if not, accept it with a probability \exp(-1/2*((x^2)new - (x^2)old))
    log_accept_prob = -1/2*(new_chisq - chisq)
    # make it a log to avoid computing exponentials
    if np.log(np.random.rand(1)) < log_accept_prob:</pre>
        return new params, new chisq
    else:
        return params, chisq
# main mcmc function
def mcmc main(d,initial params,cov,errs,nstep=20000):
   Main function for the MCMC chain
    # initialize chain
    n = initial params.size
    chain_params = np.zeros((nstep,n),dtype=float)
    chain params[0,:] = initial params
    # compute initial chi squared
    pred = get spectrum(params=initial params, take params=True)[:len(d)]
    initial chisq = chisquared(d,pred,errs)
    chain chisq = np.zeros(nstep,dtype=float)
    chain chisq[0] = initial chisq
    # take `nstep` number of steps
    for i in range(1,nstep):
        # get the old parameters
        params = chain params[i-1,:]
        chisq = chain chisq[i-1]
        # compute the putatively new ones
        params , chisq = mcmc step(d,params,chisq,cov,errs)
        # put them into the chain
        chain_params[i,:] = params_
        chain chisq[i] = chisq
    return chain params, chain chisq
# code to run on terminal in camb environment
if name == ' main ':
    # import data
    planck = np.loadtxt('COM PowerSpect CMB-TT-full R3.01.txt',skiprows=1)
    ell = planck[:,0]
    spec = planck[:,1]
    errs = (planck[:,2] + planck[:,3]) / 2
    # initial quess
    p0 = np.asarray([69, 0.022, 0.12, 0.054, 2.1e-9, 0.95])
```

```
# get model spectrum
pred = get_spectrum(params=p0,take_params=True)
pred = pred[:len(spec)]
params_newton = np.loadtxt('planck_fit_params.txt')[:,0] # for curvature me
grad = params_grad(fun=get_spectrum,params=params_newton)
grad = grad[:len(spec)]
# covariance matrix
Ninv = np.linalg.inv(np.diag(errs**2))
lhs = grad.T@Ninv@grad
cov = np.linalg.inv(lhs)
# run the mcmc simulation
nsteps = 10000 # takes O(1) second per step, so this should take ~3 hours
t1 = time.time()
chain_params, chain_chisq = mcmc_main(d=spec,initial_params=p0,cov=cov,errs
t2 = time.time()
# save the data
np.savetxt("planck chain tau.txt", chain params)
np.savetxt("planck_chisq_tau.txt", chain_chisq)
# report time
print("Time it took to do {} MCMC steps:{}".format(nsteps,t2-t1))
```

The resulting chains are saved in the files:

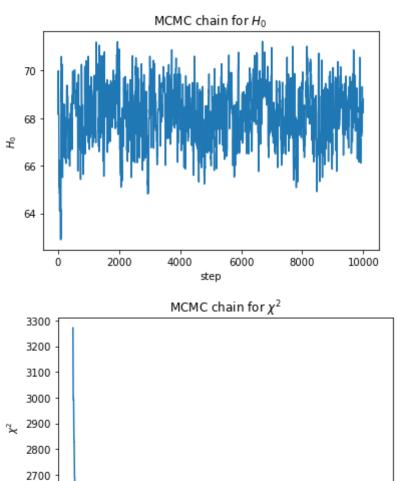
```
In [242... planck_chain = np.loadtxt("mcmc/planck_chain.txt")
    planck_chisq = np.loadtxt("mcmc/planck_chisq.txt")
```

Which, at first glance, yield the putative convergences:

```
In [246... x = np.arange(10000)

plt.figure()
plt.title(r"MCMC chain for $H_0$")
plt.ylabel("$H_0$")
plt.xlabel("step")
plt.plot(x,planck_chain[:,0])
plt.show()

plt.figure()
plt.title(r"MCMC chain for $\chi^2$")
plt.ylabel("$\chi^2$")
plt.xlabel("step")
plt.xlabel("step")
plt.plot(x,planck_chisq)
plt.axhline(2501,c='k',ls='--')
plt.show()
```



So, did the values converge? Hard to say. Judging by the upper plot of H_0 as a function of MCMC steps, I would say that no, the MCMC algorithm has not converged. Retracing back our steps, I think this is due to a bug in my original Newton's method iterator (which is odd, since this code worked fine for all other problem sets and I still cannot seem to find the bug after many attempts and building a minimal model to test it -- however, since the get_spectrum function is more or less a black box, testing our grad_params function on it is not very enlightening). This in turn yields a faulty covariance matrix which trickles down into our MCMC algorithm, providing dubious new test parameters. I have done my best to attempt to remedy this by tweaking the step with an overall scale factor of order unity (as was done in problem set 4) and varying the way in which steps are taken (e.g. trying out a uniform distribution) but the results are qualitatively the same. The trial and error method proved at least somewhat effective, as it yielded a $<\chi^2>pprox n=2501$, but this method is time-intensive, since every trial takes an order of six hours. So, I do not think my chains are converged, and I can trace back the cause to the erroneous covariance matrix obtained via Newton's method stemming from a bug (which I have not discovered due to the black box nature of get_spectrum not liking grad_params though not without much toil) when <code>grad_params</code> acts on <code>get_spectrum</code>.

2600 2500

0

2000

4000

6000

step

8000

10000

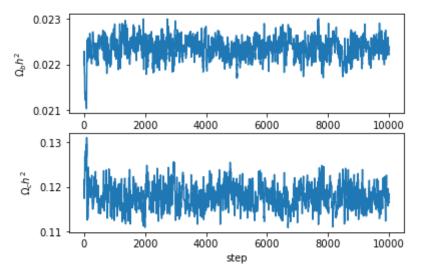
Moving on as though everything had worked, what are the values and their errors?

```
In [244...
         # set a burn-in time
         burnin = 100 # more or less
         # these are the best parameters (again, assuming convergence)
         mcmc params = planck chain[-1,:]
         # the error is simply the standard deviation of the sample
         sigma_mcmc_params = np.std(planck_chain[burnin:,:],axis=0)
         print("H0 = {} +/- {}".format(mcmc params[0], sigma mcmc params[0]))
         print("Omegabh2 = {} +/- {}".format(mcmc_params[1],sigma_mcmc_params[1]))
         print("Omegach2 = {} +/- {}".format(mcmc_params[2],sigma_mcmc_params[2]))
         print("tau = {} +/- {}".format(mcmc_params[3],sigma_mcmc_params[3]))
         print("As = {} +/- {}".format(mcmc_params[4],sigma_mcmc_params[4]))
         print("ns = {} +/- {}".format(mcmc_params[5],sigma_mcmc_params[5]))
         H0 = 68.80375944904863 + - 1.0872354015286563
         Omegabh2 = 0.02221693293836467 +/- 0.00020751209881565068
         Omegach2 = 0.11650370781229581 +/- 0.002436431723053146
         tau = 0.09546878602718682 +/- 0.02604136802420618
         As = 2.264039671665958e - 09 + / - 1.0923033350069017e - 10
         ns = 0.9745173040304714 +/- 0.005852441622004641
```

Again, I'm fairly certain this isn't correct. Indeed, the error in some values (H_0 and τ) is on the same order as the first significant figure. Also, the value of τ is significantly off from our initial guess and the value presented in question 4). One redeeming feature is that $\Omega_{b,c}h^2$ seem to have a reasonably good (samll) error with respect to their value. Looking at their MCMC chains, we can see that their convergence is once again not complete:

```
In [245... fig,ax=plt.subplots(nrows=2)
   plt.suptitle(r"MCMC chain for $\Omega$'s'")
   ax[0].set_ylabel("$\Omega_b h^2$")
   ax[1].set_ylabel("$\Omega_c h^2$")
   ax[1].set_xlabel("step")
   ax[0].plot(x,planck_chain[:,1])
   ax[1].plot(x,planck_chain[:,2])
   plt.show()
```

MCMC chain for Ω's'



In the interest of completing this assignment, however, we move on yet. The value of h is extracted from H_0 as such:

```
In [162... h=planck_chain[-1,0]/100
```

We may now solve for Ω_{Λ} using $1=\Omega_{\Lambda}+\Omega_b+\Omega_c$. We use a mini monte-carlo (not mcmc) method to propagate errors (i.e. create a bunch of Gaussian distributed samples centred at the mean with deviation equal to the error, and get the final error by computing the standard deviation of the total).

```
In [189... # set up the means and standard deviations
         lots = 10000
         H0 = mcmc_params[0]
         H0 sigma = sigma mcmc params[0]
         omega_bh = mcmc_params[1]
         omega_bh_sigma = sigma_mcmc_params[1]
         omega_ch = mcmc_params[2]
         omega_ch_sigma = sigma_mcmc_params[2]
         # set up the normally distributed arrays
         H0s = np.random.normal(loc=H0, scale=H0_sigma, size=lots)
         omega_bhs = np.random.normal(loc=omega_bh, scale=omega_bh_sigma, size=lots)
         omega_chs = np.random.normal(loc=omega_ch, scale=omega_ch_sigma, size=lots)
         # calculate everything
         hs = H0s / 100
         omega_lambdas = 1 - omega_bhs / hs**2 - omega_bhs / hs**2
         omega lambda = np.mean(omega lambdas)
         omega lambda sigma = np.std(omega lambdas)
         # print it
         print("Omega Lambda is: {} +/- {}".format(omega lambda,omega lambda sigma))
         Omega_Lambda is: 0.906105480381642 +/- 0.003079592729882093
```

So we have obtained $\Omega_{\Lambda}=0.906(3).$

4)

We will now repeat the MCMC procedure of the last problem, however with the additional constraint that $\tau=0.054\pm0.0074$. We include this constraint simply by adding a term in our χ^2 of the form <code>chisq_tau = ((trial_tau-tau_prior))/(sigma_tau_prior))**2</code> where <code>tau_prior +/- sigma_tau_prior = 0.054 \pm 0.0074</code>. This can be thought of as an additional model constraint (we have an updated prior, if you will). So, let's run a new chain with this constraint:

```
In [236... planck_chain_tau = np.loadtxt("mcmc/planck_chain_tau.txt")
    planck_chisq_tau = np.loadtxt("mcmc/planck_chisq_tau.txt")

# the params are the last site
    mcmc_params_tau = planck_chain_tau[-1,:]
```

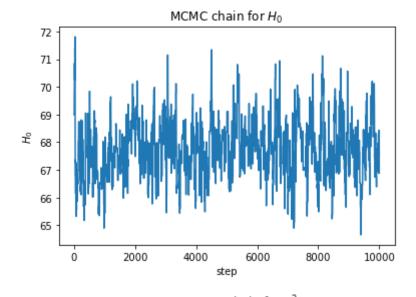
```
# the errors are the standard deviation of the chain
sigma_mcmc_params_tau = np.std(planck_chain_tau[burnin:,:], axis=0)
```

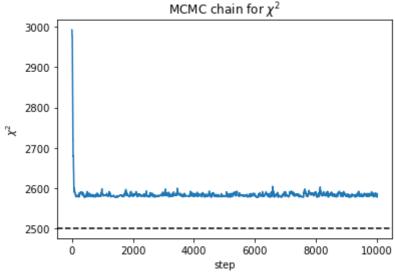
Let's have a quick look at convergence:

```
In [240... x = np.arange(10000)

plt.figure()
plt.title(r"MCMC chain for $H_0$")
plt.plot(x,planck_chain_tau[:,0])
plt.ylabel("$H_0$")
plt.xlabel("step")
plt.show()

plt.figure()
plt.title(r"MCMC chain for $\chi^2$")
plt.ylabel("$\chi^2$")
plt.ylabel("$\chi^2$")
plt.xlabel("step")
plt.xlabel("step")
plt.xlabel("step")
plt.plot(x,planck_chisq_tau)
plt.axhline(2501,c='k',ls='--')
plt.show()
```





The same problems as before are inherited here as well. These are the parameters we

obtain:

```
In [237... print("H0 = {} +/- {}".format(mcmc_params_tau[0], sigma_mcmc_params_tau[0]))
    print("Omegabh2 = {} +/- {}".format(mcmc_params_tau[1], sigma_mcmc_params_tau[1]
    print("Omegach2 = {} +/- {}".format(mcmc_params_tau[2], sigma_mcmc_params_tau[2]
    print("tau = {} +/- {}".format(mcmc_params_tau[3], sigma_mcmc_params_tau[3]))
    print("As = {} +/- {}".format(mcmc_params_tau[4], sigma_mcmc_params_tau[4]))
    print("ns = {} +/- {}".format(mcmc_params_tau[5], sigma_mcmc_params_tau[5]))

H0 = 67.61733091750979 +/- 1.0394051189437972
    Omegabh2 = 0.02229257109929826 +/- 0.00021165372624995315
    Omegach2 = 0.11934489549145502 +/- 0.002360128663603689
    tau = 0.05415361478650203 +/- 0.007222597642066387
    As = 2.0933171769615404e-09 +/- 3.148098100992451e-11
    ns = 0.9688233530956443 +/- 0.0060204364942864795
```

Next we want to do importance sampling on the chain of the last problem with the additional $\delta\chi^2$ term (stemming from the contraint of τ) acting as the effective weight and compare our results to those obtained by running a whole new chain. The code is as follows:

```
In [ ]: # import data
        planck = np.loadtxt('COM PowerSpect CMB-TT-full R3.01.txt', skiprows=1)
        ell = planck[:,0]
        spec = planck[:,1]
        errs = (planck[:,2] + planck[:,3]) / 2
        # import chains
        planck chain = np.loadtxt("planck chain.txt")
        planck chisq = np.loadtxt("planck chisq.txt")
        # define the new parameter value
        tau prior = 0.0540
        sigma tau prior = 0.0074
        # compute the new chi2 (effective weight)
        # i.e. just the difference in our old taus and the new constrained one
        tau_chain = planck_chain[:,3]
        planck chisq new = ((tau prior-tau chain)/sigma tau prior)**2
        # with this new chisquared, compute the new phase-space density L'/L
        density = np.exp(-0.5 * (planck chisq new))
        # compute the new parameters as weighted sums
        density all = np.tile(density,(planck chain.shape[1],1)).T / np.sum(density)
        params_weighted = np.sum(density_all * planck_chain, axis=0)
        # save the data
        np.savetxt("planck chisq new.txt", planck chisq new)
        np.savetxt("planck density.txt", density)
        np.savetxt("planck params weighted.txt", params weighted)
```

Running this, we obtain the new values for the parameters. We should also re-calculate the errors by taking the deviation of our new weighted chains.

```
In [233... planck_chain_weighted = np.loadtxt("mcmc/planck_chain_weighted.txt")
    sigma_mcmc_params_sampling = np.std(planck_chain_weighted[burnin:,:], axis=0)
```

```
In [234... | mcmc_params_sampling = np.loadtxt("mcmc/planck_params_weighted.txt")

print("H0 = {} +/- {}".format(mcmc_params_sampling[0],sigma_mcmc_params_sampling | print("Omegabh2 = {} +/- {}".format(mcmc_params_sampling[1],sigma_mcmc_params_s | print("Omegach2 = {} +/- {}".format(mcmc_params_sampling[2],sigma_mcmc_params_s | print("tau = {} +/- {}".format(mcmc_params_sampling[3],sigma_mcmc_params_sampling | print("As = {} +/- {}".format(mcmc_params_sampling[4],sigma_mcmc_params_sampling | print("ns = {} +/- {}".format(mcmc_params_sampling[5],sigma_mcmc_params_sampling | mcmc_params_sampling | mcmc_params_sampl
```

The values of H_0 , $\Omega_b h^2$, $\Omega_c h^2$, and A_s are consistent with running the entire chain again with a modified χ^2 to account for the τ constraint. The values of τ and n_s are not, however.