

tomcarron__exercise10

June 20, 2022

1 Exercise Set 10

Due: **9:30 20 June 2022**

Discussion: **13:00 24 June 2022**

Online submission at via [ILIAS](#) in the directory Exercises / Übungen -> Submission of Exercises / Rückgabe des Übungsblätter

```
[ ]: import numpy as np
import pandas as pd
import astropy
import astropy.constants as const
import astropy.units as u
import matplotlib.pyplot as plt
import matplotlib
import seaborn as sb
import scipy
import scipy.integrate
import math
matplotlib.rcParams['figure.dpi']=300
matplotlib.rcParams['figure.figsize']=(4,3)
```

2 1. Maximum Likelihood analysis [100 points]

2.1 Flux density measurements

In this exercise we repeat an example from the lecture and extend it: **Flux density measurements** (Lecture Note 11 p. 17).

Like in the lecture we suppose that we have flux density measurements at the frequencies,

$$f_i \in (0.4, 1.4, 2.7, 5, 10) \text{ GHz},$$

with corresponding measured flux densities,

$$1.855, 0.640, 0.444, 0.22, 0.102 \text{ flux units.}$$

Let the frequencies be f_i and the data S_i . The measurements follow a power-law slope of -1 , on top of which a 10% Gaussian noise level is added. The noise level is denoted ϵ and the model for the flux density as a function of frequency is $k f^{-\gamma}$. Assuming we know the noise level and distribution, each term in the likelihood product is of the form:

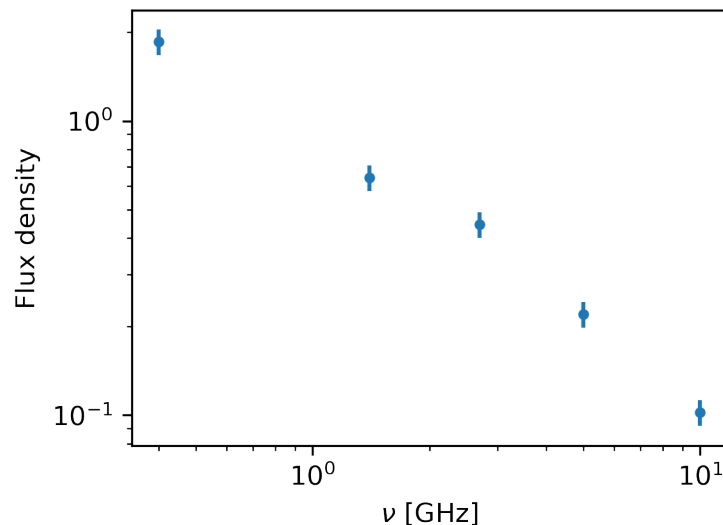
$$\frac{1}{\sqrt{2\pi}\epsilon k f_i^{-\gamma}} \exp\left(-\frac{(S_i - k f_i^{-\gamma})^2}{2(\epsilon k f_i^{-\gamma})^2}\right)$$

We are therefore fitting Gaussians to the errors.

```
[ ]: #frequencies and fluxes
fi=np.array([0.4,1.4,2.7,5,10]) #Giga-Hertz
si=np.array([1.855, 0.640, 0.444, 0.22, 0.102]) #flux units
alpha=-1 #spectral index
```

a) Plot the measured spectrum on a log-log scale. **10 points**

```
[ ]: plt.errorbar(fi,si,si*0.1,fmt='. ')
plt.xlabel(r"$\nu$ [GHz]")
plt.ylabel(r"Flux density")
plt.yscale("log")
plt.xscale("log")
plt.savefig("plots/fig_a.png",dpi=300,bbox_inches="tight")
```



b) Compute the logarithmic likelihood for a range of k and γ values in the range of the maximum likelihood. Normalize the likelihood. Produce a 2-D plot of the logarithmic likelihood for each calculated pair of k and γ . **20 points**

```

[ ]: N=2000 #Number of ks and gammas
kmin,kmax=0.1,2
ks=np.linspace(kmin,kmax,N)
gammas=np.linspace(kmin,kmax,N)
noise=0.1

#Function to compute the log likelihood function for a given k,gamma and noise_
↪level.
def log_like_prod(k,gamma,si,fi,epsilon):
    sum=0.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        #print(i,"/n",pre,"/n",exp_term)
        sum+=(exp_term-np.log(pre))
    return sum

def like_prod(k,gamma,si,fi,epsilon):
    prod=1.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        prod*=(np.exp(exp_term)/pre)
    return prod

#Function re-ordered for integrating over gamma
def like_prod2(gamma,k,si,fi,epsilon):
    prod=1.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        prod*=(np.exp(exp_term)/pre)
    return prod

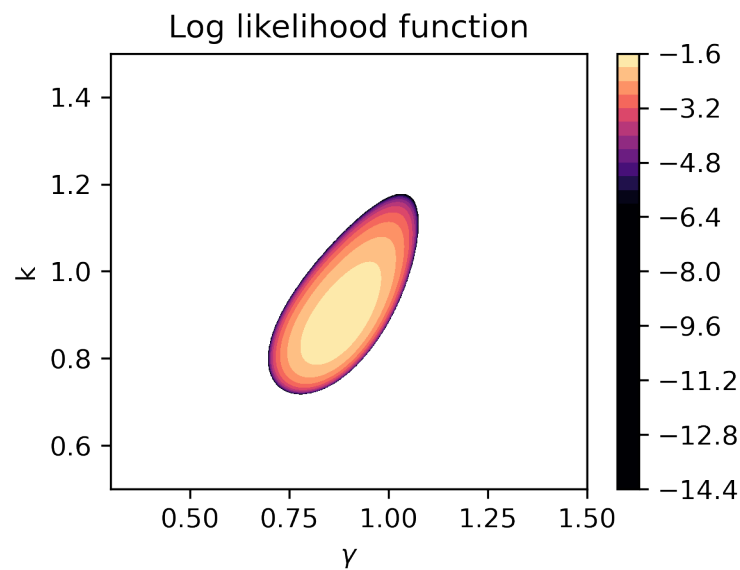
#normalization
norm,norm_err=scipy.integrate.dblquad(like_prod, kmin,kmax,kmin,kmax,
    ↪args=(si,fi,noise))

```

```
[ ]: x,y=(np.meshgrid(ks,gammas))
result=log_like_prod(x,y,si,fi,noise)
z=np.log(result/norm)

plt.contourf(y,x,z,cmap='magma',levels=30,vmin=-6)
plt.xlabel(r"$\gamma$")
plt.ylabel(r"$k$")
plt.colorbar()
plt.xlim(0.3,1.5)
plt.ylim(0.5,1.5)
#plt.yscale("log")
#plt.xscale("log")
plt.title("Log likelihood function")
plt.savefig("plots/fig_b.png",dpi=300,bbox_inches="tight")
```

<ipython-input-6-d855408caa53>:3: RuntimeWarning: invalid value encountered in log
 z=np.log(result/norm)



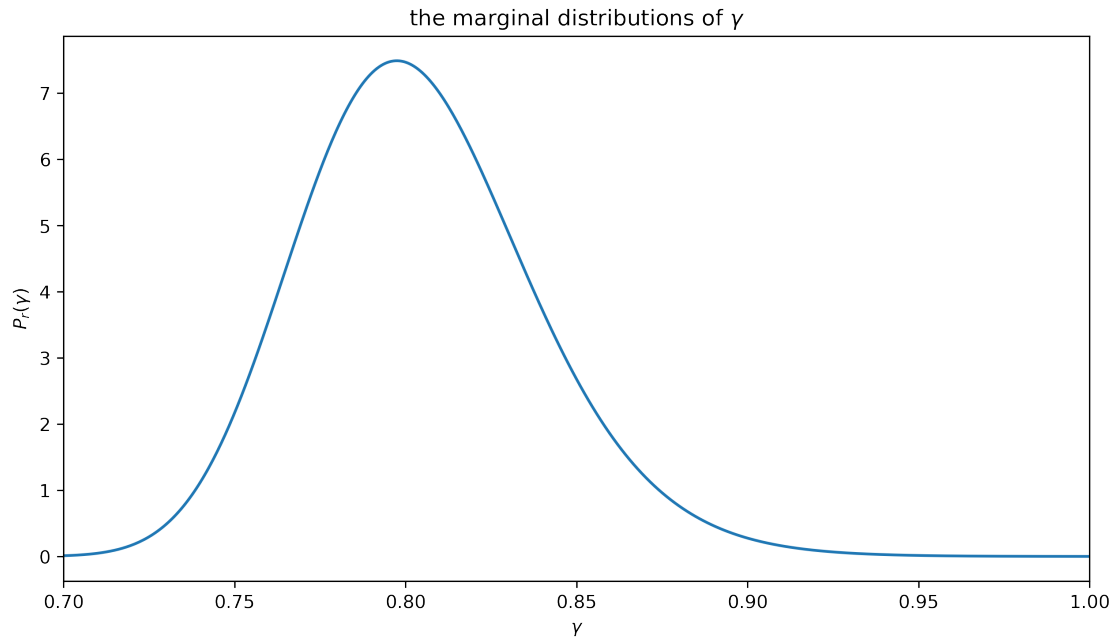
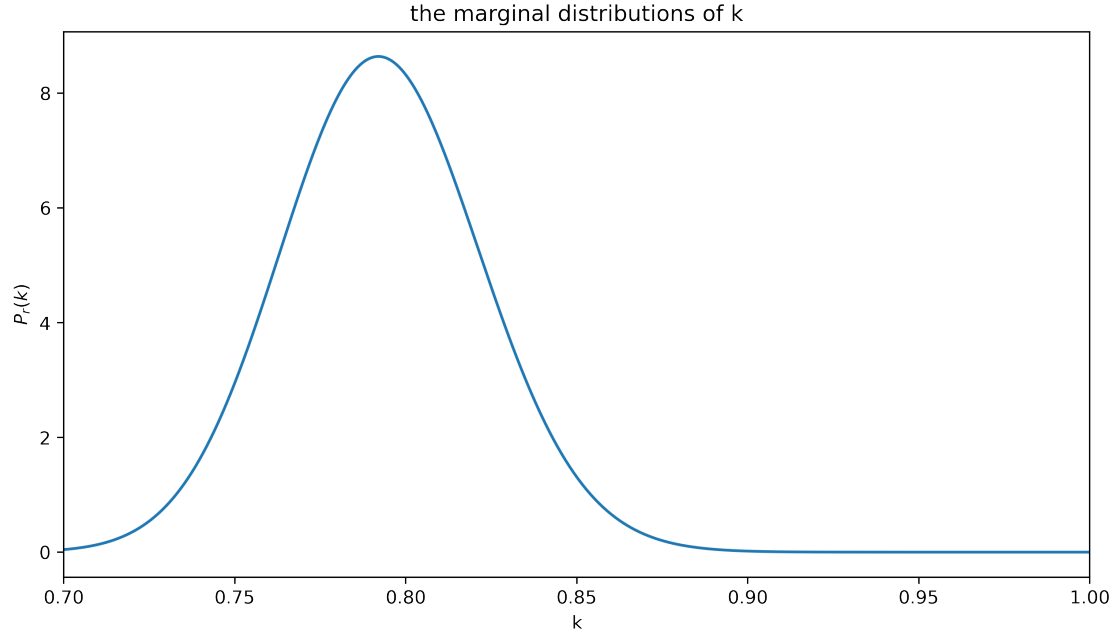
c) Now look at the probability of one parameter independent of the other parameter. This is called marginalization. In other words we have the posterior probability $\mathcal{P}_r(k, \gamma | S_i)$ and can form:

$$\mathcal{P}_r(k | S_i) = \int \mathcal{P}_r(k, \gamma | S_i) d\gamma$$

Compute the marginal probability distributions for k and γ and plot them individually. **20 points**

```
[ ]: #marginal probability of k
k_mp=[]
for i in range(len(gammas)):
    gamma=gammas[i]
    k_mp.append(scipy.integrate.quad(like_prod,0.3,1.
    ↪5,args=(gamma,si,fi,noise))[0]/norm)
#marginal probability of gamma
gamma_mp=[]
for i in range(len(ks)):
    k=ks[i]
    gamma_mp.append(scipy.integrate.quad(like_prod2,0.3,1.
    ↪5,args=(k,si,fi,noise))[0]/norm )
```

```
[ ]: ks2=np.linspace(0.3,1.5,len(ks))
gammas2=np.linspace(0.3,1.5,len(gammas))
fig, axs = plt.subplots(2, 1,figsize=(10, 12))
axs[0].plot(ks2,k_mp)
axs[0].set_xlabel(r"k")
axs[0].set_ylabel(r"$P_r(k)$")
axs[1].plot(gammas2,gamma_mp)
axs[1].set_xlabel(r"$\gamma$")
axs[1].set_ylabel(r"$P_r(\gamma)$")
axs[0].set_title(r"the marginal distributions of k")
axs[1].set_title(r"the marginal distributions of $\gamma$")
axs[0].set_xlim(0.7,1.0)
axs[1].set_xlim(0.7,1.0)
plt.savefig("plots/marginal.png",dpi=300,bbox_inches="tight")
```



d) Add an *artificial* 0.4 flux units to the flux measurements, then use it to calculate two models:

- Model A is the same as above with no offsets.
- Model B uses a model for the flux densities of the form of $\beta + kf^{-\gamma}$. Thus each likelihood term is:

$$\frac{1}{\sqrt{2\pi}\epsilon kf_i^{-\gamma}} \exp\left(-\frac{(S_i - (\beta + kf_i^{-\gamma}))^2}{2(\epsilon kf_i^{-\gamma})^2}\right)$$

Plot the contours of the computed logarithmic likelihoods for each Model into the same plot. Compare both models with each other. Which one gives better results? Explain why. **30 points**

```
[ ]: def model_b(beta,k,gamma,si,fi,epsilon):
    sum=0.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=beta+(k*(f**(-1.0*gamma)))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        sum+=-(exp_term-np.log(pre))
    return sum

def like_prod_b(k,gamma,si,fi,epsilon,beta):
    prod=1.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=beta+k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        prod*=(np.exp(exp_term)/pre)
    return prod

def like_prod_b2(gamma,k,si,fi,epsilon,beta):
    prod=1.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=beta+k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        prod*=(np.exp(exp_term)/pre)
    return prod

def like_prod_b3(beta,gamma,k,si,fi,epsilon):
    prod=1.0
    for i in range(len(si)):
        s=si[i]
        f=fi[i]
        temp=beta+k*(f**(-1.0*gamma))
        pre=((np.sqrt(2*np.pi)*epsilon*temp))
        exp_term=((-(s-temp)**2)/(2*(epsilon*temp)**2))
        prod*=(np.exp(exp_term)/pre)
    return prod
```

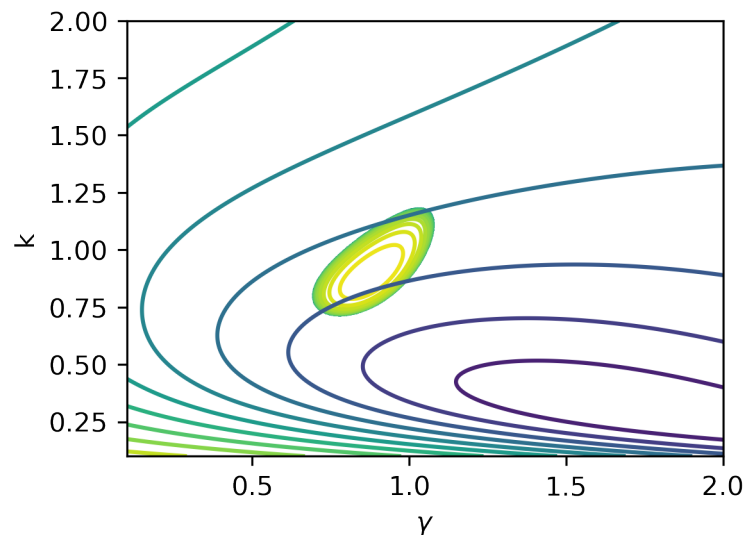
```
norm2,norm_err2=scipy.integrate.dblquad(like_prod_b, kmin,kmax,kmin,kmax,
↪args=(si,fi,noise,0.4))
```

```
[ ]: x,y=(np.meshgrid(ks,gammas))
result=log_like_prod(x,y,si,fi,noise)
z=np.log(result/norm)

result2=(model_b(0.4,x,y,si,fi,noise))
z2=np.log(result2/norm2)

plt.contour(y,x,z,levels=30)
plt.contour(y,x,z2,levels=10)
plt.xlabel(r"$\gamma$")
plt.ylabel(r"$k$")
#plt.xlim(0.3,1.5)
#plt.ylim(0.5,1.5)
plt.savefig("plots/fig_modelb.png",dpi=300,bbox_inches="tight")
```

<ipython-input-10-ab85afd47593>:3: RuntimeWarning: invalid value encountered in log
 z=np.log(result/norm)



e) Compute the marginal probability distributions for Model B for the three parameters: β , γ , and k and plot them. **20 points**

```
[ ]: #marginal probability of k for model b
k_mp=[]
for i in range(len(gammas)):
    gamma=gammas[i]
```



```

    k_mp.append(scipy.integrate.quad(like_prod_b,0.0,5.
↪0,args=(gamma,si,fi,noise,0.4))[0]/norm2 )
#marginal probability of gamma model b
gamma_mp=[]
for i in range(len(ks)):
    k=ks[i]
    gamma_mp.append(scipy.integrate.quad(like_prod_b2,0.0,5.
↪0,args=(k,si,fi,noise,0.4))[0]/norm2 )

#marginal probability of beta model b
#:/

```

```

[ ]: ks2=np.linspace(0.0,5.0,len(ks))
    gammas2=np.linspace(0.0,5.0,len(gammas))
    fig, axs = plt.subplots(2, 1,figsize=(10, 12))
    axs[0].plot(ks2,k_mp)
    axs[0].set_xlabel(r"k")
    axs[0].set_ylabel(r"$P_r(k)$")
    axs[1].plot(gammas2,gamma_mp)
    axs[1].set_xlabel(r"$\gamma$")
    axs[1].set_ylabel(r"$P_r(\gamma)$")
    axs[0].set_title(r"the marginal distributions of k")
    axs[1].set_title(r"the marginal distributions of $\gamma$")
    #axs[0].set_xlim(0.7,1.0)
    #axs[1].set_xlim(0.7,1.0)
    plt.savefig("plots/marginal2.png",dpi=300,bbox_inches="tight")

```

