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### CS20BTECH11063

### **Data Science Analysis Assignment 7**

```
import numpy as np
import matplotlib.pyplot as plt
import scipy.stats as stats
from scipy import optimize
import astroML
from astroML.stats import sigmaG
import pandas as pd
import seaborn as sns
np.random.seed(0)
import emcee
import corner
from sklearn.neighbors import KernelDensity
from IPython.display import display, Math
import dynesty
import nestle
```

Q1

$$f_{gas}=f_0(1+f_1z)$$

```
In [14]: df = pd.read_csv('q1.txt', sep=' ')
    z = df['z']
    fgas = df['fgas']
    fgas_error = df['fgas_error']

def log_prior(theta):
    f0, f1 = theta
    if 0 < f0 < 0.5 and -0.5 < f1 < 0.5:
        return 0.0
    return -np.inf</pre>
```

```
def log likelihood(theta, x, y, sigma y):
    f0, f1 = theta
    model = f0 * (1 + f1 * x)
    chi2 = np.sum(np.log(2 * np.pi * sigma y ** 2) + ((y - model) ** 2 / sigma y ** 2))
    return -0.5 * chi2
def log likelihood2(theta, x, y, sigma y):
    f0, f1 = theta
    model = f0 * (1 + f1 * x)
    chi2 = np.sum(np.log(2 * np.pi * sigma y ** 2) + ((y - model) ** 2 / sigma y ** 2))
    return 0.5 * chi2
def log probability(theta, x, y, sigma y):
    lp = log prior(theta)
    if not np.isfinite(lp):
        return -np.inf
    return lp + log likelihood(theta, x, y, sigma y)
# Solution from scipy.optimize.minimize
optimized sol = optimize.fmin(log likelihood2, [0.2, 0.0], args=(z, fgas, fgas error))
# # plot the results
# f0, f1 = optimized sol
# print(f0, f1)
\# x = np.linspace(0, 1.5, 1000)
# y = f0 * (1 + f1 * x)
# plt.figure(figsize=(15, 5))
# plt.errorbar(z, fgas, yerr=fgas_error, fmt='.', color='black', capsize=5, ecolor='gray', label='Data')
# plt.plot(x, y, color='red')
# plt.xlabel('z')
# plt.ylabel('fgas')
# plt.ylim(0.06, 0.21)
# plt.xlim(0, 1.5)
# plt.grid()
# plt.show()
# Solution from emcee
ndim, nwalkers = 2, 64
# f0 random = np.random.uniform(0, 0.5, nwalkers)
# f1 random = np.random.uniform(-0.5, 0.5, nwalkers)
\# pos = np.array([f0_random, f1_random]).T \#optimized_sol.x + 1e-4 \# np.random.randn(nwalkers, ndim)
pos = optimized sol + 1e-4 * np.random.randn(nwalkers, ndim)
sampler = emcee.EnsembleSampler(nwalkers, ndim, log probability, args=(z, fgas, fgas error))
nburn = 1000
nsteps = 2000
```

```
sampler.run mcmc(pos, nsteps, progress=True)
samples = sampler.get chain(flat=True, discard=nburn)
# print(samples.shape)
f0 median, f1 median = np.median(samples, axis=0)
f0 68, f1 68 = np.percentile(samples, [16, 84], axis=0)
f0 90, f1 90 = np.percentile(samples, [5, 95], axis=0)
# print the results using math display
display(Math(r'f 0 = {0:.3f} +{1:.3f} -{2:.3f}) (68\% CI)'.format(f0 median, f0 68[1] - f0 median, f0 median - f0 68[0])))
display(Math(r'f 1 = \{0:.3f\} + \{1:.3f\} - \{2:.3f\}  (68\% CI)'.format(f1 median, f1 68[1] - f1 median, f1 median - f1 68[0])))
display(Math(r'f 0 = \{0:.3f\} + \{1:.3f\} - \{2:.3f\} (90\% CI)'.format(f0 median, f0 90[1] - f0 median, f0 median - f0 90[0])))
display(Math(r'f 1 = \{0:.3f\} + \{1:.3f\} - \{2:.3f\} (90\% CI)'.format(f1 median, f1 90[1] - f1 median, f1 median - f1 90[0])))
# print("f0 = {0:.3f} + {1:.3f} - {2:.3f}) (68% CI)".format(f0 median, f0 68[1] - f0 median, f0 median - f0 68[0]))
# print("f1 = \{0:.3f\} +\{1:.3f\} -\{2:.3f\} (68% CI)".format(f1 median, f1 68[1] - f1 median, f1 median - f1 68[0]))
# print("f0 = {0:.3f} + {1:.3f} - {2:.3f} (90\% CI)".format(f0 median, f0 90[1] - f0 median, f0 median - f0 90[0]))
# print("f1 = \{0:.3f\} + \{1:.3f\} - \{2:.3f\} (90\% CI)".format(f1 median, f1 90[1] - f1 median, f1 median - f1 90[0]))
# Corner plot for 68% and 90% confidence intervals
# fig = plt.figure(figsize=(10, 10))
labels = [r"$f 0$", r"$f 1$"]
fig = corner.corner(samples, labels=labels, levels=[0.68, 0.9], show titles=True, title kwargs={"fontsize": 12})
plt.show()
# plot the best fit line and the data
samples reshaped = samples.reshape((-1, ndim)).T
# print(samples reshaped.shape)
x = np.linspace(0, 1.5, 1000)
f0, f1 = samples reshaped[:2]
y = f0[:, None] * (1 + f1[:, None] * x)
# print(y.shape)
mu = np.mean(y, axis=0)
sig = 2 * y.std(0)
upper bound = mu + sig
lower bound = mu - sig
plt.figure(figsize=(15, 5))
plt.plot(x, mu, color='black', label='Best fit line')
plt.errorbar(z, fgas, yerr=fgas error, fmt='.', color='black', capsize=5, ecolor='gray', label='Data')
plt.fill between(x, lower bound, upper bound, color='grey', alpha=0.2, label='Confidence Interval Region')
plt.xlim(0, 1.5)
plt.ylim(0.06, 0.21)
plt.grid()
plt.legend()
plt.show()
```

Optimization terminated successfully.

Current function value: -200.732528

Iterations: 60

Function evaluations: 114

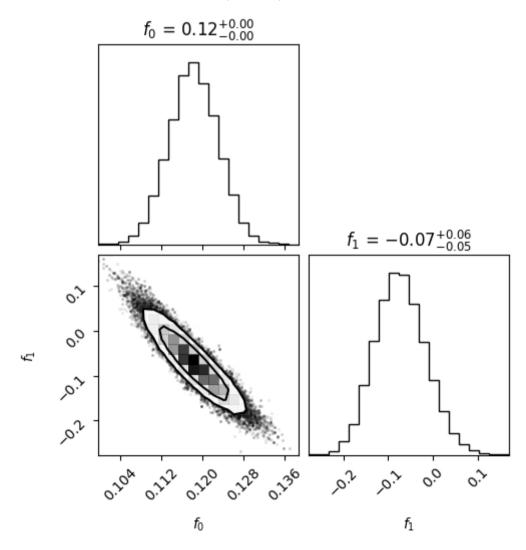
#### 100%| 2000/2000 [01:53<00:00, 17.63it/s]

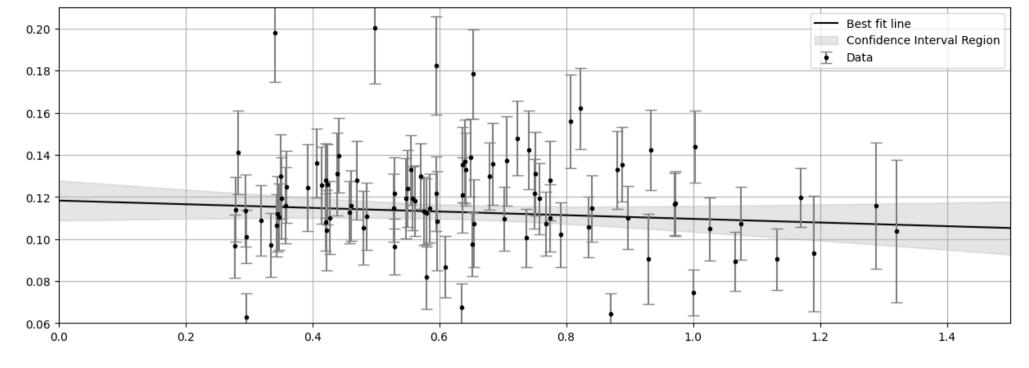
$$f_0 = 0.118 + -0.246 - 0.005(68\%CI)$$

$$f_1 = -0.074 + 0.059 - -0.197(68\%CI)$$

$$f_0 = 0.118 + -0.279 - 0.008(90\%CI)$$

$$f_1 = -0.074 + 0.098 - -0.200(90\%CI)$$





Reference: http://jakevdp.github.io/blog/2014/06/14/frequentism-and-bayesianism-4-bayesian-in-python/

## Q2

```
import numpy as np
In [17]:
        data = np.array([[ 0.42,  0.72,  0. ,  0.3 ,  0.15,
                         0.09, 0.19, 0.35, 0.4, 0.54,
                         0.42, 0.69, 0.2, 0.88, 0.03,
                         0.67, 0.42, 0.56, 0.14, 0.2 ],
                       [0.33, 0.41, -0.22, 0.01, -0.05,
                        -0.05, -0.12, 0.26, 0.29, 0.39,
                         0.31, 0.42, -0.01, 0.58, -0.2,
                         0.52, 0.15, 0.32, -0.13, -0.09],
                       [ 0.1 , 0.1 , 0.1 , 0.1 , 0.1 ,
                         0.1, 0.1, 0.1, 0.1, 0.1,
                         0.1, 0.1, 0.1, 0.1, 0.1,
                         0.1, 0.1, 0.1, 0.1, 0.1]])
        x, y, sigma_y = data
        # def polynomial_fit(theta, x):
              val = 0
             for i in range(len(theta)):
```

```
val += theta[i] * x**i
      return val
def polynomial fit(theta, x):
    """Polynomial model of degree (len(theta) - 1)"""
    return sum(t * x ** n for (n, t) in enumerate(theta))
def log likelihood(theta, x=x, y=y, sigma y=sigma y):
    model = polynomial fit(theta, x)
    # return -0.5 * np.sum((y - model)**2 / sigma y**2)
    return -0.5 * np.sum(np.log(2 * np.pi * sigma y ** 2) + (y - model) ** 2 / sigma y ** 2)
# def prior transform(theta):
      return theta
def prior transform linear(theta):
    m = theta[0]*10 - 5
    c = theta[1]*20 - 10
    return np.array([m, c])
def prior transform quadratic(theta):
    a = theta[0]*10 - 5
    b = theta[1]*10 - 5
    c = theta[2]*20 - 20
    return np.array([a, b, c])
nlive = 1024
bound = 'multi'
ndims = 2
sample = 'unif'
tol = 0.1
sampler linear = dynesty.NestedSampler(log likelihood, prior transform linear, ndims, nlive=nlive, bound=bound, sample=sample)
samples linear = sampler linear.run nested(dlogz=tol, print progress=True)
results_linear = sampler_linear.results
log_z_linear = results_linear.logz[-1]
print(results linear.summary())
nlive = 1024
bound = 'multi'
ndims = 3
sample = 'unif'
tol = 0.1
sampler_quadratic = dynesty.NestedSampler(log_likelihood, prior_transform_quadratic, ndims, nlive=nlive, bound=bound, sample=sample)
samples quadratic = sampler quadratic.run nested(dlogz=tol, print progress=True)
```

```
results quadratic = sampler quadratic.results
log z quadratic = results quadratic.logz[-1]
print(results quadratic.summary())
print("Bayes Factor: ", np.exp(log z quadratic - log z linear))
print("Bayes Factor: ", np.exp(log z linear - log z quadratic))
12187it [00:12, 944.62it/s, +1024 | bound: 10 | nc: 1 | ncall: 51471 | eff(%): 26.188 | loglstar: -inf < 22.011 <
                                                                                                                     inf | logz: 12.463
+/- 0.091 | dlogz: 0.000 > 0.100]
Summary
======
nlive: 1024
niter: 12187
ncall: 50447
eff(%): 26.188
logz: 12.463 +/- 0.117
None
14883it [00:16, 878.84it/s, +1024 | bound: 16 | nc: 1 | ncall: 62651 | eff(%): 25.812 | loglstar: -inf < 22.940 <
                                                                                                                     inf | logz: 10.753
+/- 0.104 | dlogz: 0.000 > 0.100]
Summary
======
nlive: 1024
niter: 14883
ncall: 61627
eff(%): 25.812
logz: 10.753 +/- 0.131
None
Bayes Factor: 0.18076547443894297
Bayes Factor: 5.532029847534681
```

## Q3

```
In [3]: df = pd.read_csv('q3.csv', sep='\t')
    z = df['z']
# Plot KDE estimate for z using gaussian and exponential kernels (with bandwidth=0.2)

plt.figure(figsize=(15, 5))
plt.xticks(np.arange(-1, 6, 0.5))
plt.yticks(np.arange(0, 1.1, 0.05))

x = np.linspace(-0.5, 5.5, 1000)
kde = KernelDensity(kernel='gaussian', bandwidth=0.2).fit(z.to_numpy().reshape(-1, 1))
```

```
log_dens = kde.score_samples(x.reshape(-1,1))

plt.plot(x, np.exp(log_dens), label='Gaussian Kernel (0.2)')
kde = KernelDensity(kernel='exponential', bandwidth=0.2).fit(z.to_numpy().reshape(-1, 1))
log_dens = kde.score_samples(x.reshape(-1,1))
plt.plot(x, np.exp(log_dens), label='Exponential Kernel (0.2)')
plt.hist(z, bins='auto', density=True, color='grey', alpha=0.5, label='Data')
plt.xlabel('z')
plt.ylabel('Probability Density')
plt.legend()
plt.grid()
plt.show()
```

