```
In [1]: import matplotlib.pyplot as plt
import numpy as np
import pandas as pd
from scipy.stats import cauchy
from scipy.stats import norm
import seaborn as sn
import korali
from scipy.optimize import fsolve
from IPython.display import display, HTML
display(HTML("<style>.container { width:85% !important; }</style>"))
import random
```

AM207 Homework 4 - Bayesian inference

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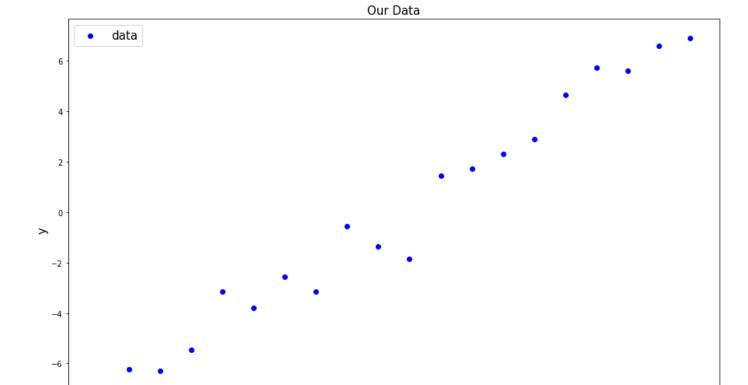
Date: 11/17/2022

Question 1: Bayesian inference of linear model

An experimentalist collected data $\{x_i,y_i\}_{i=1}^N$. A model of this data is given by $y_i=ax_i+\sigma\xi_i$ where a and σ are scalars and ξ_i are i.i.d. normally distributed random variables, $\xi_i\sim\mathcal{N}(0,1)$. We denote our parameters as $\theta=[a,\sigma]^T\in\mathbb{R}^2$. We define $d_i:=y_i-ax_i=\xi_i$ which is a random variable following $d_i\sim\mathcal{N}(0,\sigma^2)$. Indeed, $(d_i)_{1\leq i\leq N}$ are independent as for the random variables ξ_i .

```
In [2]: df_1 = pd.read_csv('./1_laplace/data.csv')
    x = df_1["x"].to_numpy()
    Y = df_1["y"].to_numpy()
    fig = plt.figure(figsize = (15,10))

plt.scatter(X,Y, c = 'b', label = 'data')
    plt.xlabel('x', fontsize = 15)
    plt.ylabel('y', fontsize = 15)
    plt.title('Our Data', fontsize = 15)
    plt.legend(loc = 'upper left', fontsize = 15)
    plt.show()
```



a)

-8

Let $i \in \{1, \cdots, N\}$. We have that the likelihood of each d_i given parameters $heta = (a, \sigma)$ is :

$$p(d_i \mid heta) = rac{1}{\sqrt{2\pi\sigma^2}} \mathrm{exp}igg(-rac{(y_i - ax_i)^2}{2\sigma^2}igg)$$

Thus the likelihood of d_1, \dots, d_n is the product of each likelihood $p(d_i \mid \theta)$ since we assume each response d_i to be independent. Hence, we have that the likelihood of the data is :

$$p(D \mid heta) = \prod_{i=1}^N p(d_i \mid heta) = \prod_{i=1}^N rac{1}{\sqrt{2\pi\sigma^2}} \mathrm{exp}igg(-rac{(y_i - ax_i)^2}{2\sigma^2}igg)$$

where we define by D our data $(\{x_i\}_{1 \leq i \leq N}, \{y_i\}_{1 \leq i \leq N})$.

Finally, we have that the log-likelihood of the data is:

$$= \log \left(\prod_{i=1}^{N} p(d_i \mid \theta) \right)$$

$$= \sum_{i=1}^{N} \log[p(d_i \mid \theta)]$$

$$= \sum_{i=1}^{N} \log \left[\frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{(y_i - ax_i)^2}{2\sigma^2}\right) \right]$$

$$= N \log \left[\frac{1}{\sqrt{2\pi\sigma^2}} \right] - \sum_{i=1}^{N} \frac{(y_i - ax_i)^2}{2\sigma^2}$$

$$= -\frac{N}{2} \log(2\pi\sigma^2) - \sum_{i=1}^{N} \frac{(y_i - ax_i)^2}{2\sigma^2}$$

$$= -\frac{N}{2} \log(2\pi) - N \log(\sigma) - \frac{1}{2\sigma^2} \sum_{i=1}^{N} (y_i - ax_i)^2$$
(1)

Let us now find the parameters $\theta_0=(a_0,\sigma_0)$ that maximise the log-likelihood. To do so, let's differentiate the log likelihood function with respect to (a,σ) and find the roots of the loglikelihood's derivative.

 $\mathcal{L} = \log(p(D \mid \theta))$

$$0 = \frac{\partial \mathcal{L}}{\partial a} \mid_{\theta_0} = -\frac{1}{2\sigma_0^2} \sum_{i=1}^N (2a_0 x_i^2 - 2y_i x_i) \implies a_0 = \frac{\sum_{i=1}^N x_i y_i}{\sum_{i=1}^N x_i^2}$$

$$0 = \frac{\partial \mathcal{L}}{\partial \sigma} \mid_{\theta_0} = -\frac{N}{\sigma_0} + \frac{1}{\sigma_0^3} \sum_{i=1}^N (y_i - ax_i)^2$$

$$\implies \sigma_0^2 = \frac{1}{N} \sum_{i=1}^N (y_i - a_0 x_i)^2 \text{ since } \sigma_0 \neq 0$$

$$\implies \sigma_0^2 = \frac{1}{N} \sum_{i=1}^N \left(y_i - x_i \frac{\sum_{j=1}^N x_j y_j}{\sum_{j=1}^N x_j^2} \right)^2$$

$$(2)$$

In order to check whether or not the likelihood is maximal at (a_0, σ_0) , we must check that the Hessian of the log likelihood is negative definite.

$$\frac{\partial^{2} \mathcal{L}}{\partial a^{2}} \Big|_{\theta_{0}} = -\frac{1}{\sigma_{0}^{2}} \sum_{i=1}^{N} x_{i}^{2} \leq 0$$

$$\frac{\partial^{2} \mathcal{L}}{\partial \sigma^{2}} \Big|_{\theta_{0}} = \frac{N}{\sigma_{0}^{2}} - \frac{3}{\sigma_{0}^{4}} \sum_{i=1}^{N} (y_{i} - a_{0}x_{i})^{2} = \frac{N}{\sigma_{0}^{2}} - \frac{3N}{\sigma_{0}^{2}} = \frac{-2N}{\sigma_{0}^{2}} \leq 0$$

$$\frac{\partial^{2} \mathcal{L}}{\partial a \partial \sigma} \Big|_{\theta_{0}} = \frac{-2}{\sigma_{0}^{3}} \sum_{i=1}^{N} (y_{i}x_{i} - a_{0}x_{i}^{2}) = 0$$

$$\frac{\partial^{2} \mathcal{L}}{\partial \sigma \partial a} \Big|_{\theta_{0}} = \frac{2}{\sigma_{0}^{3}} \sum_{i=1}^{N} (a_{0}x_{i}^{2} - y_{i}x_{i}) = 0$$
(3)

Hence the hessian ${f H}$ of the log-likelihood evaluated at $heta_0$ is diagonal :

$$\mathbf{H}_{ heta_0} = \left(egin{array}{ccc} rac{\partial^2 \mathcal{L}}{\partial a^2} & rac{\partial^2 \mathcal{L}}{\partial a \partial \sigma} \ rac{\partial^2 \mathcal{L}}{\partial \sigma \partial a} & rac{\partial^2 \mathcal{L}}{\partial \sigma^2} \end{array}
ight) (heta_0) = \left(egin{array}{ccc} rac{\partial^2 \mathcal{L}}{\partial a^2} & 0 \ 0 & rac{\partial^2 \mathcal{L}}{\partial \sigma^2} \end{array}
ight) (heta_0)$$

and negative definite as we have that for all $\theta^T = [a, \sigma]^T \in \mathbb{R}^2$:

$$egin{aligned} heta^T \mathbf{H}_{ heta_0} heta &= a^2 rac{\partial^2 \mathcal{L}}{\partial a^2}(a_0, \sigma_0) + b^2 rac{\partial^2 \mathcal{L}}{\partial \sigma^2}(a_0, \sigma_0) \leq 0 \end{aligned}$$

Hence, we have that that the loglikelihood and the likelihood are maximal at the point $\theta_0 = (a_0, \sigma_0)$.

```
In [3]: N = len(X)
a_0 = np.sum(X*Y)/np.sum(X**2)
sigma_0 = np.sqrt((1/N)*np.sum((Y - a_0*X)**2))
theta_0 = np.array([a_0, sigma_0])
print(f'The point at which the likelihood and log-likelihood function is maximal is: (a
```

The point at which the likelihood and log-likelihood function is maximal is: (a,sigma) = (1.4864344460664443, 0.7795182956841904).

b)

Let us now express the posterior distribution of the parameters. We assume uniform priors with large enough bounds for a and σ , respectively a_1,a_2 and σ_1,σ_2 . We define $I:=[a_1,a_2]\times [\sigma_1,\sigma_2]\subset \mathbb{R}^2$. As our parameters follow a uniform distribution, their pointwise density is a constant which we denote as $p(\theta):=\frac{1}{C}\in\mathbb{R}$.

By Baye's Theorem, we have that:

$$p(\theta \mid D) = \frac{p(D \mid \theta)p(\theta)}{p(D)}$$

$$= \frac{p(D \mid \theta)}{Cp(D)}$$

$$= \frac{p(D \mid \theta)}{C\int_{I} p(D \mid \theta)p(\theta)d\theta}$$

$$= \frac{p(D \mid \theta)}{C\int_{I} p(D \mid \theta)\frac{1}{C}d\theta}$$

$$= \frac{p(D \mid \theta)}{\int_{I} p(D \mid \theta)d\theta}$$

$$\propto p(D \mid \theta) \text{ since } \int_{I} p(D \mid \theta)d\theta \text{ is a constant}$$

$$(4)$$

The log-Laplace approximation for our log posterior distribution around the equilibria $heta_0=(\sigma_0,a_0)^T$ is:

$$egin{aligned} \log p(heta \mid D) &\propto \mathcal{L}(D \mid heta_0) +
abla_{ heta} \mathcal{L}(D \mid heta_0)^T (heta - heta_0) + rac{1}{2} (heta - heta_0)^T \mathbf{H}_{ heta_0} (heta - heta_0) \end{aligned} \ &pprox \mathcal{L}(D \mid heta_0) + rac{1}{2} (heta - heta_0)^T \mathbf{H}_{ heta_0} (heta - heta_0) ext{ since the log-likelihood is maximal at } heta_0 \end{aligned} \ &= \mathcal{L}(D \mid heta_0) + rac{\partial^2 \mathcal{L}(D \mid heta)}{\partial a^2} |_{ heta_0} (a - a_0)^2 + rac{\partial^2 \mathcal{L}(D \mid heta)}{\partial \sigma^2} |_{ heta_0} (\sigma - \sigma_0)^2 \end{aligned}$$

c)

y = np.linspace(0, 1.5, 100)

We have that our target distribution is $p(\theta \mid D)$. We form the following proposal distributions from two independent Cauchy distributions with median at the previous sample value and with a scale parameter $\gamma:Q_1(\sigma_1,\sigma_0,\gamma)=\frac{1}{\pi\gamma}\left[\frac{\gamma^2}{(\sigma_1-\sigma_0)^2+\gamma^2}\right]$, $Q_2(a_1,a_0,\gamma)=\frac{1}{\pi\gamma}\left[\frac{\gamma^2}{(a_1-a_0)^2+\gamma^2}\right]$. We note that since the proposal cauchy distributions are symmetric, the acceptance rate becomes: $p(\theta_{proposed}\mid D)/p(\theta_{current}\mid D)$

```
In [4]: def likelihood(theta):
             'returns likelihood of the data D = (X, Y) given parameters theta'
            a = theta[0]
            sigma = theta[1]
            coef = 1/((2*np.pi*(sigma**2))**(N/2))
            S = np.sum((Y - a*X)**2)
            S \exp = np.exp(-S/(2*sigma**2))
            return coef*S exp
In [5]: ###first derivatives
        partial a theta 0 = \text{np.sum}(X*Y - a 0*X**2) ## this is 0
        partial s theta 0 = (-N/sigma\ 0 + np.sum((Y - a\ 0*X)**2)/(sigma\ 0**3)) ## this is 0
        ###double derivatives
        partial_aa_theta_0 = -1/(sigma_0**2)*np.sum(X**2)
        partial ss theta 0 = -2*N/sigma 0**2
        print(f'log likelihood double derivative with respect to a at theta 0 is : {partial aa t
        log likelihood double derivative with respect to a at theta 0 is : -303.15296336969317
        log likelihood double derivative with respect to sigma at theta 0 is : -65.8275006174190
In [6]:
        def posterior(theta):
             'returns posterior up to the normalisation constant'
            return likelihood(theta)
        def log posterior(theta):
            'returns the log of the exact posterior'
            return np.log(posterior(theta))
        def posterior log laplace(theta):
             'returns the log-laplace approximation of log-posterior density'
            a = theta[0]
            sigma = theta[1]
            return likelihood(theta 0) + partial aa theta 0 * (a - a 0)**2 + partial ss theta 0
        def posterior laplace(theta):
            'returns the exponential of the log laplace approxmation of the log posterior'
            return np.exp(posterior log laplace(theta))
In [7]: fig, (ax1, ax2) = plt.subplots(1, 2, subplot kw=dict(projection='3d'), figsize = (20,20)
        x = np.linspace(1, 2, 100)
```

```
X \text{ val}, Y \text{ val} = \text{np.meshgrid}(x, y)
Z = posterior laplace([X val, Y val])
ax1.contour3D(X val, Y val, Z, 50, cmap='viridis') #cmap = binary, viridis
ax1.scatter([a 0],[sigma 0], [posterior laplace(theta 0)], s = 70, c = 'r', label = 'The
ax1.legend()
ax1.set xlabel('a', fontsize = 15)
ax1.set ylabel('$\sigma$', fontsize = 15)
ax1.set zlabel(r'\$hat{p}(\lambda b)', fontsize = 15);
ax1.set title('Laplace Approximation of Posterior distribution around maximum', fontsize
x = np.linspace(0., 3, 100)
y = np.linspace(-2.5, 4, 100)
X \text{ val}, Y \text{ val} = \text{np.meshgrid}(x, y)
Z = posterior log laplace([X val, Y val])
ax2.contour3D(X val, Y val, Z, 50, cmap='viridis') #cmap = binary, viridis
ax2.scatter([a 0],[sigma 0], [posterior log laplace(theta 0)], s = 70, c = 'r', label =
ax2.legend()
ax2.set xlabel('a', fontsize = 15)
ax2.set_ylabel('$\sigma$', fontsize = 15)
ax2.set zlabel(r'\ \log \hat{p}(\theta \mid D\)', fontsize = 15);
ax2.set title('Log-Laplace Approximation of Log-Posterior distribution around maximum',
plt.show()
Laplace Approximation of Posterior distribution around maximum
                                                   Log-Laplace Approximation of Log-Posterior distribution around maximum

    Theorical maximum

    Theorical maximum

                                           1.0
                                           0.6
                                                                                                 6000
                                          θ)
0.4 <sup>©</sup>
                                          0.2
```

-1200

-1400

where we denote in the z axis \hat{p} as the laplace approximation of the posterior.

✓ 1.0 0.8

0.6

0.4

1.2

```
In [8]: def MH(max_iter, gamma, theta):
    '''x represents a; y represents sigma; this function implements Metropolis Hasting to sample parameters (a, sigma) from the log posterior density'''

t, x, y = 0, theta[0], theta[1]
Sx, Sy = [], []
p = log_posterior(np.array([x, y]))
accept, reject, dx, dy = 0, 0, 0, 0
Ra, Rs = [], []

while True:
    Sx.append(x)
    Sy.append(y)
    t += 1
```

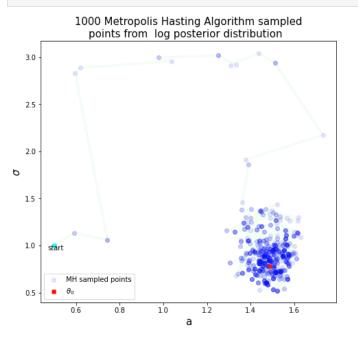
0.0

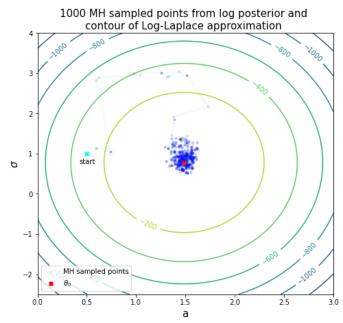
1.0

```
break
                 dx = cauchy.rvs(scale = gamma)
                 dy = cauchy.rvs(scale = gamma)
                 xp, yp = x + dx, y + dy
                 thetap = np.array([xp,yp])
                 pp = log posterior(thetap)
                 u = np.random.uniform(low=0., high=1.0, size = 1)[0]
                 if (pp > p \text{ or } pp >= np.log(u) + p) and y > 0 and yp > 0: # y,yp > 0 to enforce
                     x, y = xp, yp
                     p = pp
                      accept += 1
                 else:
                     Ra.append(xp)
                     Rs.append(yp)
                     reject += 1
              print(f"Rejection rate for \u03B3 = \{gamma\} is : ", reject/(accept + reject))
              return np.array(Sx), np.array(Sy), np.array(Ra), np.array(Rs), reject/(accept + reje
In [9]: gamma = 0.1
         max iter= 1000
         theta start = [0.5, 1]
         print('We chose gamma in order to have a rejection rate close to 0.7: \n')
         Sx1, Sy1, Ra1, Rs1, R1 = MH (max iter, gamma, theta start)
         We chose gamma in order to have a rejection rate close to 0.7:
         Rejection rate for \gamma = 0.1 is : 0.6996996996997
         /Users/elieattias/anaconda3/lib/python3.7/site-packages/ipykernel launcher.py:7: Runtime
         Warning: divide by zero encountered in log
           import sys
In [10]: fig, (ax1, ax2) = plt.subplots(1, 2)
         fig.set figheight(7)
         fig.set figwidth(17)
         ax1.plot(Sx1, Sy1, alpha=0.1, c = 'lightgreen', lw=3)
         ax1.scatter(Sx1, Sy1, alpha=0.1, c = 'b', label = 'MH sampled points')
         ax1.scatter([a 0], [sigma 0], c = 'r', label = r'$\theta 0$', marker = 'X')
         ax1.scatter([theta start[0]], [theta start[1]], c = 'cyan', marker = 'X')
         ax1.text(x = theta start[0] - 0.03, y = theta start[1] - 0.05, s = 'start')
         ax1.legend(loc = 'lower left', fontsize = 10)
         ax1.set xlabel('a', fontsize = 15)
         ax1.set ylabel(f'$\sigma$', fontsize = 15)
         ax1.set title(f' 1000 Metropolis Hasting Algorithm sampled \npoints from log posterior
         ## contour plot
         x = np.arange(np.min(Sx1) - 0.5, np.max(Sx1) + 0.5, 0.1)
         y = np.arange(np.min(Sy1) - 0.5, np.max(Sy1) + 0.5, 0.1)
         x = np.linspace(0., 3, 100)
         y = np.linspace(-2.5, 4, 100)
         x \log = np.linspace(0., 3, 100)
         y log = np.linspace(-2.5, 4, 100)
         X \text{ plot}, Y \text{ plot} = \text{np.meshgrid}(x, y)
         X plot, Y plot = np.meshgrid(x log, y log)
         Z plot = np.array([[posterior log laplace(np.array([X plot[i,j],Y plot[i,j]])) for j in
         CS = plt.contour(X plot, Y plot, Z plot)
         ax2.plot(Sx1, Sy1, alpha=0.1, c = 'lightgreen', lw=3)
         ax2.scatter(Sx1, Sy1, alpha=0.1, c = 'b', label = 'MH sampled points', s = 10)
```

if t>=max iter:

```
ax2.scatter([a_0], [sigma_0], c = 'r', label = r'$\theta_0$', marker = 'X')
ax2.scatter([theta_start[0]], [theta_start[1]], c = 'cyan', marker = 'X')
ax2.text(x = theta_start[0] - 0.08, y = theta_start[1] - 0.25, s = 'start')
ax2.clabel(CS, inline=1, fontsize=10)
ax2.legend(loc='lower left', fontsize = 10)
ax2.legend(loc='lower left', fontsize = 10)
ax2.set_xlabel('a', fontsize=15)
ax2.set_ylabel('$\sigma$', fontsize=15)
ax2.set_title('1000 MH sampled points from log posterior and \ncontour of Log-Laplace applit.show()
```





d)

Now we have K=1000 samples points $\{a_k,\sigma_k\}$. For each sample pair (a_k,σ_k) , we can hence make k predictions on y given x. A prediction y given a sample item $\theta_k=(a_k,\sigma_k)$ follows a normal distribution i.e $y \mid x \sim \mathcal{N}(a_k x,\sigma_k^2)$ where $k \in \{1,\cdots,1000\}$.

Hence, for any x,y, we have 1000 predictions for $p(y\mid D,x)$. We approximate such prediction by the mean :

$$p(y \mid D; x) pprox rac{1}{K} \sum_{k=1}^K rac{1}{\sqrt{2\pi\sigma_k^2}} \mathrm{exp}igg(-rac{(y-a_k x)^2}{2\sigma_k^2}igg)$$

We then approxmiate the cdf by the cumulative sum with respect to y:

$$C(y) = \sum_{i=min(Y)}^{y} p(i \mid D; x)$$

Finally, to find the quantile y_q , we compute the following:

$$y_q = \operatorname{argmin}_y \left(|C(y) - q| \right)$$

These are shown for x=-4 and x=3 in the examples here-under.

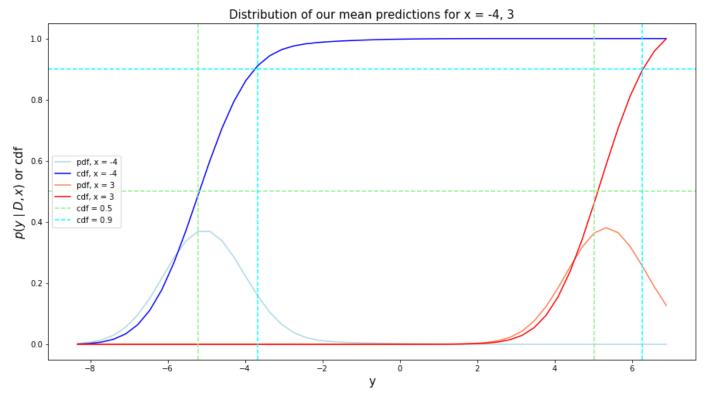
```
In [11]: L = 50 K = 1000
```

```
y_val = np.linspace(np.min(Y), np.max(Y), L)
x_val = np.linspace(np.min(X), np.max(X), L)
```

```
In [12]: x1 = -4
    pdfs_2 = np.array([[norm.pdf(y_val[k] - x1* Sx1[i], Sy1[i]) for k in range(L)] for i in
    pdf_mean_2 = np.mean(pdfs_2, axis = 0)
    cdf_2 = np.cumsum(pdf_mean_2)
    cdf_2 = cdf_2/np.max(cdf_2)
    q1 = y_val[np.argmin(np.abs(cdf_2 - 0.5))]
    q2 = y_val[np.argmin(np.abs(cdf_2 - 0.9))]

x2 = 3
    pdfs_3 = np.array([[norm.pdf(y_val[k] - x2* Sx1[i], Sy1[i]) for k in range(L)] for i in
    pdf_mean_3 = np.mean(pdfs_3, axis = 0)
    cdf_3 = np.cumsum(pdf_mean_3)
    cdf_3 = cdf_3/np.max(cdf_3)
    q3 = y_val[np.argmin(np.abs(cdf_3 - 0.5))]
    q4 = y_val[np.argmin(np.abs(cdf_3 - 0.9))]
```

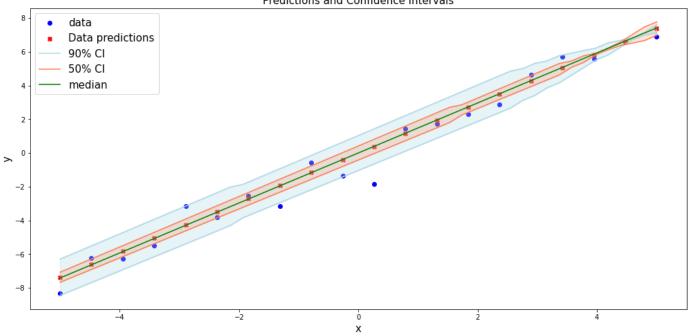
```
In [13]: fig = plt.figure(figsize = (15,8))
         plt.plot(y val, pdf mean 2, c ='lightblue', label= f'pdf, x = {x1}')
         plt.plot(y val, cdf 2/np.max(cdf 2), c = 'b', label = f'cdf, x = {x1}')
         plt.plot(y val, pdf mean 3, c = 'coral', label= f'pdf, x = {x2}')
         plt.plot(y val, cdf 3/np.max(cdf 3), c = 'r', label = f'cdf, x = {x2}')
         plt.axhline(0.5, c = 'lightgreen', linestyle = 'dashed', label = 'cdf = 0.5')
         plt.axhline(0.9, c = 'cyan', linestyle = 'dashed', label = 'cdf = 0.9')
         plt.axvline(q1, c = 'lightgreen', linestyle = 'dashed')
         plt.axvline(q2, c = 'cyan', linestyle = 'dashed')
         plt.axvline(q3, c = 'lightgreen', linestyle = 'dashed')
         plt.axvline(q4, c = 'cyan', linestyle = 'dashed')
         plt.xlabel('y', fontsize = 15, c = 'black')
         plt.ylabel(r'p(y \in D, x) or cdf', fontsize = 15, c = 'black')
         plt.title(f'Distribution of our mean predictions for x = \{x1\}, \{x2\}', fontsize = 15, c =
         plt.legend(loc = 'center left')
         plt.show()
```



```
In [15]: quantiles_50 = np.zeros(L)
quantiles_90 = np.zeros(L)
```

```
medians = np.zeros(L)
         for c, x in enumerate(x val):
             predict y = x*Sx1
             predict y.sort()
             median = predict y[K//2] #since K = 1000 is even
             pdfs = np.array([[norm.pdf(y val[1] - \times \times Sx1[k], Sy1[k]) for 1 in range(L)] for k i.
             pdf mean = np.mean(pdfs, axis = 0)
             cdf = np.cumsum(pdf mean)
             cdf = cdf/np.max(cdf)
             y = 50 = y \text{ val}[np.argmin(np.abs(cdf - 0.5))] #we can take the argmin of the abs because
             y 90 = y val[np.argmin(np.abs(cdf - 0.9))]
             medians[c] = median
              quantiles 50[c] = y 50
              quantiles 90[c] = y 90
In [16]: predictions = x val*np.mean(Sx1)
         data predictions = X*np.mean(Sx1)
          r 50 = np.abs(predictions - quantiles 50)
          r 90 = np.abs(predictions - quantiles 90)
         y_{low_50} = predictions - r_{50/2}
         y high 50 = predictions + r 50/2
         y low 90 = predictions - r 90/2
         y high 90 = predictions + r 90/2
In [17]: fig = plt.figure(figsize = (17,8))
         plt.scatter(X, Y, c = 'b', label = 'data')
         plt.scatter(X, data predictions, c = 'r', label = 'Data predictions', marker = 'X')
         plt.fill_between(x_val, y_low_90, y_high_90, color = 'lightblue', alpha = 0.3)
         plt.fill between(x val, y low 50, y high 50, color = 'coral', alpha = 0.2)
         plt.plot(x val, y low 90, c = 'lightblue', label = '90% CI')
         plt.plot(x val, y high 90, c = 'lightblue')
         plt.plot(x val, y low 50, c = 'coral', label = '')
         plt.plot(x val, y high 50, c = 'coral', label = '50% CI')
         plt.plot(x val, medians, c = 'g' , label = 'median')
         plt.legend(loc = 'upper left', fontsize = 15)
         plt.xlabel('x', fontsize= 15)
         plt.ylabel('y', fontsize= 15)
         plt.title('Predictions and Confidence Intervals', fontsize= 15)
         plt.show()
```





Question 2: Where is the beacon?

A beacon positioned at unknown location (a,b) emits light in random directions $\theta \sim \mathcal{U}(-\pi/2,\pi/2)$. The light is detected by sensors located on the shore (y=0). The light detections are recorded at coordinates x_i , $i=1,2,\ldots,N$.

a)

We define g such that $x=g(\theta):=a-b\tan\theta$. We are given that the density of θ is uniform such that $p_{\theta}(\theta_i)=\frac{1}{\pi}$. We note that g and all our random variables X,θ are continuous. In addition, g is a bijective continuous function and is hence invertible. Its inverse is $g^{-1}(x)=\arctan\left(\frac{a-x}{b}\right)$. We have by transformation of continuous random variables that the likelihood of each x given θ and parameters $\omega=(a,b)$ is :

$$p_{X}(x \mid \omega) = p_{\theta}(g^{-1}(x) \mid \omega) \left| \frac{d}{dx} g^{-1}(x) \right|$$

$$= \frac{1}{\pi} \left| \frac{d}{dx} \arctan\left(\frac{a - x}{b}\right) \right|$$

$$= \frac{1}{\pi} \left| \frac{1}{1 + \frac{(a - x)^{2}}{b^{2}}} \times \left(\frac{-1}{b}\right) \right|$$

$$= \frac{1}{\pi} \frac{b}{b^{2} + (a - x)^{2}}$$
(6)

which is a Cauchy distribution.

Thus the likelihood of x_1, \dots, x_n is the product of each likelihood $p(x_i \mid \omega)$ since we assume each response x_i to be independent. Hence, we have that the likelihood of the data is :

$$P(X\mid\omega) = \prod_{i=1}^N p_X(x_i\mid\omega) = \prod_{i=1}^N rac{1}{\pi} rac{b}{b^2 + (a-x_i)^2}$$

where X denotes $(x_i)_{1 \leq i \leq N}$.

Hence, the log likelihood is:

$$\mathcal{L}(X \mid \omega) = \log[P(X \mid \omega)] = \sum_{i=1}^N \logigg(rac{1}{\pi} rac{b}{b^2 + (a-x_i)^2}igg)$$

b)

Let us express the posterior distribution of $\omega=(a,b)$ where both a and b are unknown. We assume that the priors are independent -i.e that $P(\omega\mid X)=P(a\mid X)P(b\mid X)$. Hence the posterior distribution for ω is:

$$p(\omega \mid X) = \frac{P(X \mid \omega)P(\omega)}{P(X)}$$

$$= \frac{P(X \mid \omega)P(a \mid X)P(b \mid X)}{P(X)}$$

$$\propto P(X \mid \omega)P(a \mid X)P(b \mid X)$$
(7)

where we assume that our priors are such that $a \mid X \sim \mathcal{U}(-20, 20)$, and $b \mid X \sim \mathcal{U}(0, 20)$. Indeed, considering that the impact points are mostly distributed around 0, it makes sense to take a prior for a symmetric around 0. Then, as understood by the problem set, the beacon's y coordinate (b) is positive.

c)

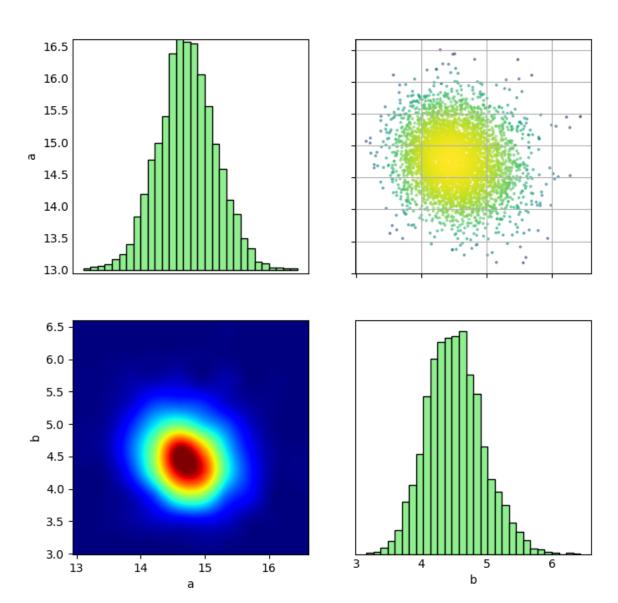
Using Korali, we sampled the posterior distribution of $\omega=(a,b)$ given the data X. The mean of the parameters a,b is a=14.76 and b=4.52. The posterior density of the beacon's position is shown here-under.

```
In [20]: df = pd.read csv('2 lighthouse/data.csv')
         x = df["x"].to numpy()
         n = len(x)
         def log likelihood(ks):
             a,b = ks["Parameters"]
             log like = np.sum(np.log(p x(a,b)))
             ks["logLikelihood"] = log like
         def p x(a,b):
             'pointwise likelihood'
             return (1/np.pi) * (b/(b**2 + (a - x)**2))
         e = korali.Experiment()
         e["Problem"]["Type"] = "Bayesian/Custom"
         e["Problem"]["Likelihood Model"] = log likelihood
         e["Solver"]["Type"] = "Sampler/TMCMC"
         e["Solver"]["Population Size"] = 5000
         e["Solver"]["Target Coefficient Of Variation"] = 1.0
         e["Solver"]["Covariance Scaling"] = 0.2
```

```
e["Distributions"] = [
   {"Name": "Prior a",
       "Type": "Univariate/Uniform",
       "Minimum": -20,
       "Maximum": 20},
   {"Name": "Prior b",
       "Type": "Univariate/Uniform",
       "Minimum": 0,
       "Maximum": 20}
e["Variables"] = [
    {"Name": "a", "Prior Distribution": "Prior a"},
   {"Name": "b", "Prior Distribution": "Prior b"}
]
e["Store Sample Information"] = True
e["Console Output"]["Verbosity"] = "Detailed"
k = korali.Engine()
k.run(e)
[Korali] -----
[Korali] Current Generation: #1
[Korali] Annealing Exponent:
                                 0.000e+00.
[Korali] Acceptance Rate (proposals / selections): (100.00% / 48.16%)
[Korali] Coefficient of Variation: 100.00%
[Korali] log of accumulated evidence: -8.969
[Korali] max logLikelihood: -804.029
[Korali] Number of finite Evaluations (prior / likelihood): (5000 / 5000)
[Korali] Sample Mean:
a = +1.268e+01
b = +8.889e+00
[Korali] Sample Covariance:
  | +5.336e+00 -
  | -1.383e+00 +4.396e+00 |
[Korali] Experiment: 0 - Generation Time: 4.548s
[Korali] -----
[Korali] Current Generation: #2
[Korali] Annealing Exponent:
                                  9.453e-03.
[Korali] Acceptance Rate (proposals / selections): (79.70% / 48.10%)
[Korali] Coefficient of Variation: 100.00%
[Korali] log of accumulated evidence: -29.977
[Korali] max logLikelihood: -804.030
[Korali] Number of finite Evaluations (prior / likelihood): (4511 / 4511)
[Korali] Sample Mean:
a = +1.465e+01
b = +6.131e+00
[Korali] Sample Covariance:
  | +1.037e+00 -
  | -1.353e-01 +1.375e+00 |
[Korali] Experiment: 0 - Generation Time: 4.604s
[Korali] -----
[Korali] Current Generation: #3
[Korali] Annealing Exponent:
                                  3.395e-02.
[Korali] Acceptance Rate (proposals / selections): (81.66% / 47.10%)
[Korali] Coefficient of Variation: 100.00%
[Korali] log of accumulated evidence: -84.031
[Korali] max logLikelihood: -804.030
[Korali] Number of finite Evaluations (prior / likelihood): (4892 / 4892)
[Korali] Sample Mean:
a = +1.477e+01
b = +4.932e+00
```

```
[Korali] Sample Covariance:
  | +2.862e-01 -
  | -1.859e-02 +2.877e-01 |
[Korali] Experiment: 0 - Generation Time: 4.598s
[Korali] -----
[Korali] Current Generation: #4
[Korali] Annealing Exponent:
                                 9.965e-02.
[Korali] Acceptance Rate (proposals / selections): (82.42% / 47.52%)
[Korali] Coefficient of Variation: 100.00%
[Korali] log of accumulated evidence: -264.372
[Korali] max logLikelihood: -804.030
[Korali] Number of finite Evaluations (prior / likelihood): (4998 / 4998)
[Korali] Sample Mean:
a = +1.477e+01
b = +4.590e+00
[Korali] Sample Covariance:
  +7.844e-02 -
  | -5.536e-03 +7.250e-02 |
[Korali] Experiment: 0 - Generation Time: 4.626s
[Korali] -----
[Korali] Current Generation: #5
[Korali] Annealing Exponent: 3.224e-01.
[Korali] Acceptance Rate (proposals / selections): (82.88% / 48.88%)
[Korali] Coefficient of Variation: 106.63%
[Korali] log of accumulated evidence: -810.277
[Korali] max logLikelihood: -804.027
[Korali] Number of finite Evaluations (prior / likelihood): (5000 / 5000)
[Korali] Sample Mean:
a = +1.477e+01
b = +4.497e+00
[Korali] Sample Covariance:
  | +2.472e-02 -
  | -9.305e-04 +2.203e-02 |
[Korali] Experiment: 0 - Generation Time: 4.672s
[Korali] ------
[Korali] Current Generation: #6
[Korali] Annealing Exponent:
                                 1.000e+00.
[Korali] Acceptance Rate (proposals / selections): (83.06% / 62.90%)
[Korali] Coefficient of Variation: 200.00%
[Korali] log of accumulated evidence: -810.277
[Korali] max logLikelihood: -804.027
[Korali] Number of finite Evaluations (prior / likelihood): (5000 / 5000)
[Korali] Sample Mean:
a = +1.476e+01
b = +4.520e+00
[Korali] Sample Covariance:
  | +4.214e-02 -
  | -2.790e-03 +3.844e-02 |
[Korali] Experiment: 0 - Generation Time: 4.750s
[Korali] -----
[Korali] sampler/TMCMC finished correctly.
[Korali] Termination Criterion Met: TMCMC['Target Annealing Exponent'] = 1.000000.
[Korali] Final Generation: 6
[Korali] Elapsed Time: 28.102s
```

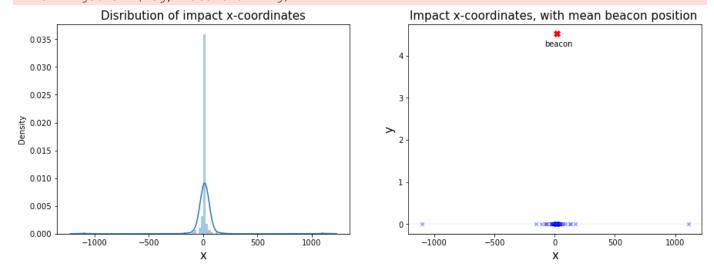
TMCMC Plotter -Number of Samples 5000



```
In [21]; fig, (ax1, ax2) = plt.subplots(1, 2, figsize = (15, 5))
         df = pd.read csv('./2 lighthouse/data.csv')
         x = df["x"].to numpy()
         sn.distplot(x,bins=100, ax = ax1)
         ax2.scatter(x,[0]*len(x), c = 'b', alpha = 0.4, s = 20, marker = 'x')
         ax1.set xlabel('x', fontsize = 15)
         ax2.set xlabel('x', fontsize = 15)
         ax2.set ylabel('y', fontsize = 15)
         a = 14.76
         b = 4.52
         ax2.scatter([a], [b], c = 'r', s = 60, marker = 'X')
         ax2.axhline(xmin = -1000, xmax = 1000, y = 0, alpha = 0.2, c = 'lightgreen')
         ax2.text(x = a-100, y = b - 0.3, s = 'beacon')
         ax1.set title('Disribution of impact x-coordinates', fontsize = 15)
         ax2.set title('Impact x-coordinates, with mean beacon position ', fontsize = 15)
         plt.show()
```

utureWarning: `distplot` is a deprecated function and will be removed in a future versio n. Please adapt your code to use either `displot` (a figure-level function with similar flexibility) or `histplot` (an axes-level function for histograms).

warnings.warn(msg, FutureWarning)



In []: