Source: https://www.youtube.com/watch?v=9_eZHt2qJs4&t=16s

KL divergence is measure how one probability distribution is different from second, reference distribution.

X is random variable and small x-s are the states it could take: $X = \{x_1, x_2, x_3, \dots, x_n\}$

We want to compare different probability distribution $\log p_{\theta}x_1$ with some other probability distribution $\log q_{\phi}x_1$. They dont have to be same type of distributions (usually q is simpler to use distribution for modelling)

One way to calculate difference is use subtraction, which because we are using logs is division:

$$\log p_{ heta}x_1 - \log q_{\phi}x_1 = \log[rac{p_{ heta}x_1}{q_{\phi}x_1}]$$

This division is called log likelihood ratio. But currently we are calculating difference between one sample. We would like to calculate average difference between p and q. In random variables we say what is expected value of the random vairable or what is its central tendency.

For random variables exptected value is weighted average of instances of random variable. Each variable and all states have probability of occurrence and average should reflect that. Samples that have higher probability should contribute more to the average.

$$\mathbb{E}_{p_{ heta}}[X] = \sum_{i=1}^{\infty} x_i p_{ heta}(x_i)$$

where:

- x_i is state of the random variable
- $p_{\theta}(x_i)$ is weight of the random variable

Here we showed that calculation should be done for very large number of samples

Here is same formula more in general, instead of random variable we calculate weighted average of a function of random variables

$$\mathbb{E}_{p_{ heta}}[h(X)] = \sum_{i=1}^{\infty} h(x_i) p_{ heta}(x_i)$$

So far we have looked at discrete random variables. For continous random variable we would calculate exptected value as such (summation has been replaced by integral):

$$\mathbb{E}_{p_{ heta}}[h(X)] = \int_{\mathbb{R}} h(x_i) p_{ heta}(x_i)$$

Our log likelihood ratio is nothig but a function of random variable and since we are interested in average of this function we should be able to use expectation. So we need weights and compute the sums:

$$\sum_{i=1}^{\infty} p_{ heta}(x_i) \log[rac{p_{ heta} x_i}{q_{\phi} x_i}]$$

To be exact we are calculating exptected value of log likelihood ratio. This is **KL divergence**. We can express it using expectation symbol:

$$\mathbb{E}_{ ext{p}}[\log[rac{p_{ heta}x_i}{q_{\phi}x_i}] = \sum_{i=i}^{\infty}p_{ heta}(x_i)\log[rac{p_{ heta}x_i}{q_{\phi}x_i}]$$

Previous example was for discrete random variable for continuous random variable we replace summation with an integral:

$$\mathbb{E}_{ ext{p}}[\log[rac{p_{ heta}x_i}{q_{\phi}x_i}] = \int_{\mathbb{R}} p_{ heta}(x_i) \log[rac{p_{ heta}x_i}{q_{\phi}x_i}]$$

We have one problem. Integral and summation both go from minus infinity to infinity. We could get help from the **law of large numbers**: "as a sample size grows, its mean gets closer to the average of the whole population" (https://www.investopedia.com/terms/l/lawoflargenumbers.asp). So we can revrite KL divergence as a mean provided we use many samples:

$$\frac{1}{N} \sum_{i=1}^{N} \log[\frac{p_{\theta} x_i}{q_{\phi} x_i}]$$

Antoher notation used is:

$$DD_{KL}(p_{ heta}||q_{\phi}) = \int_{\mathbb{R}} p_{ heta}(x) log[rac{p_{ heta}(x)}{q_{\phi}(x)}] dx$$

This on is called forward KL

If we would like to use q instead of p for weighting:

$$D_{KL}(q_{\phi}||p_{ heta}) = \int_{\mathbb{R}} q_{\phi}(x) log[rac{q_{\phi}(x)}{p_{ heta}(x)}] dx$$

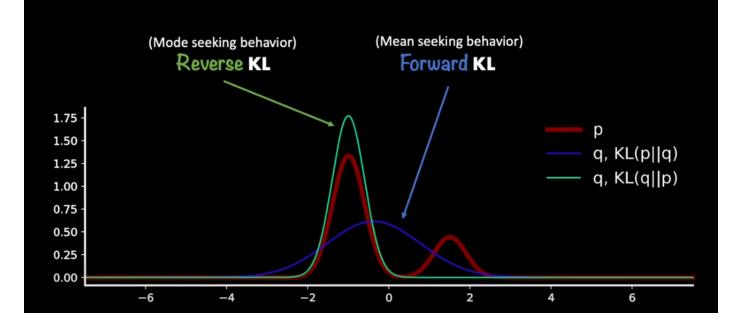
This on is called reverse KL.

These formulations are going to give different values. This is a reason why this is not called metric, but a distance.

Generally we use p for reference distribution and q for approximation.

Which one to use? It depends. Forward KL has mean-seeking behavior as reverse KL has mode seeking behavior.

```
In [89]: from IPython.display import display, Image
display(Image(filename="images/kl.png"))
```



Here we can see that reverse has picked up mean mode of the distribution. Most of the time when we are doing density estimation and using variatinal inference we use reverse scale. Forward KL is being used a lot in machine learning but you don't see name forward KL a lot. It is being used indirectly. For example when we are using cross-entropy loss in classificatin loss we are using KL indirectly (at least a component of KL divergence).

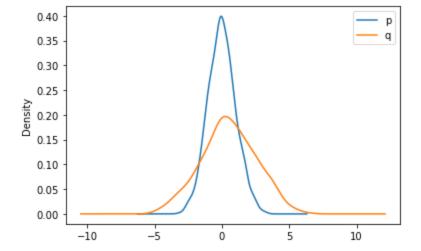
Example

```
In [1]: import pandas as pd
   import numpy as np
   from scipy.stats import norm
   import matplotlib.pyplot as plt
```

Sample KL divergence calculation

In [68]: plot_p_q_dens(samples_p, samples_q)

```
# Define the two distributions
In [65]:
         p = norm(0, 1)
         q = norm(0.5, 2)
         samples q=q.rvs(1000)
         samples p=p.rvs(1000)
         # Forward KL divergence
         forward kl = np.sum(p.logpdf(samples q) - q.logpdf(samples q))
         print("Forward KL divergence:", forward kl)
         # Reverse KL divergence
         reverse kl = np.sum(q.logpdf(samples p) - p.logpdf(samples p))
         print("Reverse KL divergence:", reverse kl)
        Forward KL divergence: -1008.6516711191375
        Reverse KL divergence: -333.72609786923385
In [66]:
        def plot p q dens(samples p, samples q):
             df q=pd.DataFrame({'q':samples q})
             df p=pd.DataFrame({'p':samples p})
             ax=df p.plot(kind='density')
             df q.plot(kind='density', ax=ax)
```



Run forward and reverse on multimodal data

Very simple example, does not estimate standard deviation, only estimates mean. Only for educational purpose not well checked or error-free

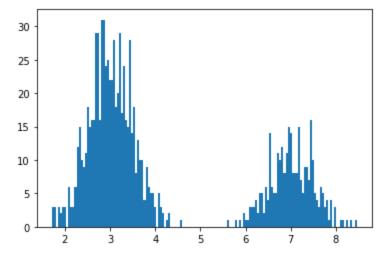
```
In [7]:
       class Pdf:
            """custom class to get pdf out of samples. Just takes proportion of data in that int
            Very simple, if value ot of sample range, takes last value probability of sample in
            def init (self, samples, n bins=100, smoothing value=0.00001):
                self.samples=samples
                self.n bins=n bins
                self.smoothing value=smoothing value
                self.bins, self.bins perc=self.create bins(self.samples, n bins=self.n bins)
                self.bins perc[self.bins perc == 0]=self.smoothing value
            def create bins(self, samples, n bins=100):
                num bin, bins=np.histogram(samples, bins=n bins)
                bins perc=num bin/sum(num bin)
                return bins, bins perc
            def get pdf(self, x):
                if isinstance(x, (np.ndarray)):
                    x=x[:,None]
                    bin idx=np.absolute((self.bins-x)).argmin(1)
                    bin idx[bin idx>len(self.bins perc)-1]=len(self.bins perc)-1
                else:
                    bin idx=np.absolute((self.bins-x)).argmin()
                    bin idx=min(bin idx, len(self.bins perc)-1)
                return self.bins perc[bin idx]
            def get logpdf(self, x):
                p=self.get pdf(x)
                return np.log(p)
            def mean(self):
                return np.mean(self.samples)
```

```
# Initialize the means and KL divergence values
means = [initial mean]
kl divergences = []
for i in range(num steps):
    if kl forward: #forward kl
        kl = np.sum(pdf p.get pdf(samples q)*(pdf p.get logpdf(samples q) - q.logpdf
        gradient = q.pdf(initial mean) * (initial mean - pdf p.mean()) / initial std
    else: #reverse kl
        kl= np.sum(q.pdf(samples)*(q.logpdf(samples) - pdf p.get logpdf(np.array(sam))
        gradient = pdf p.get pdf(initial mean) * (initial mean - pdf p.mean()) / ini
    initial mean -= step size * gradient
    kl divergences.append(kl)
    q = norm(initial mean, initial std)
    samples q=q.rvs(num samples q)
    means.append(initial mean)
plt.plot(kl divergences)
plt.xlabel('Step')
plt.ylabel('KL Divergence')
plt.show()
plt.plot(means)
plt.xlabel('Step')
plt.ylabel('Mean')
plt.show()
plot p q dens(samples, samples q)
if return data:
    return kl divergences, means, samples q
```

Generate data

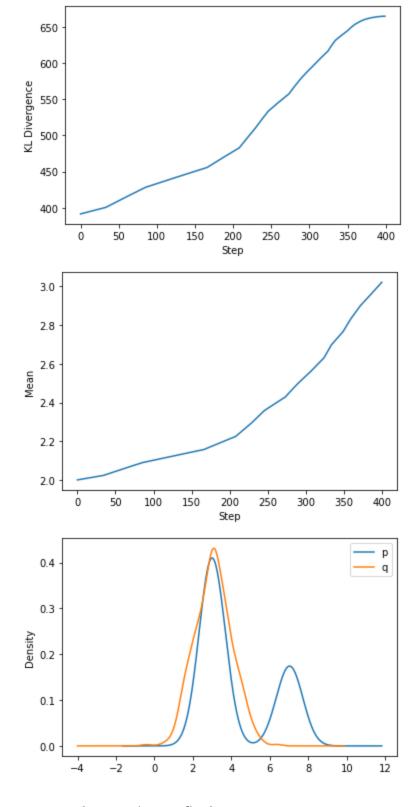
```
In [84]: #generate data we have, multimodal
    mu, sigma = 3, 0.5 # mean and standard deviation
    p_samples = np.random.normal(mu, sigma, 700)
    mu2, sigma2 = 7.0, 0.5 # mean and standard deviation
    p_samples2 = np.random.normal(mu2, sigma2, 300)
    p_samples=p_samples.tolist()+p_samples2.tolist()

    pdf_p=Pdf(p_samples)
    _=plt.hist(p_samples, bins=150)
```



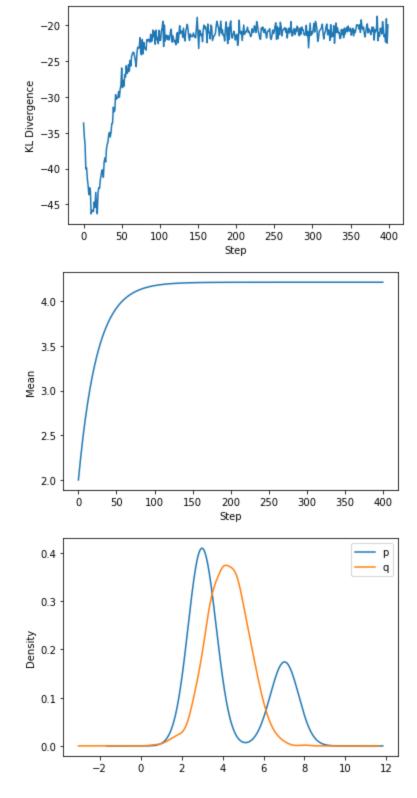
Reverse KL - tries to find one mode

In [87]: estimate_dist(p_samples, kl_forward=False, num_steps=400)



Forward KL - tries to find mean

In [91]: estimate_dist(p_samples, num_steps=400)



Convert to pdf: jupyter nbconvert --to webpdf --allow-chromium-download .\kl_divergence.ipynb