Maximum Liklehood Estimation

Mr Fugu Data Science



<u>Youtube (https://www.youtube.com/channel/UCbni-TDI-Ub8VIGaP8HLTNw/)</u> | <u>Github (https://github.com/MrFuguDataScience)</u>

Maximum Likelihood Estimation: allows us to estimate our "unknown" parameters for a given probability distribution. Think of it like this: we are trying to find the likelihood "probability" of generating this dataset.

- The parameters are found where they maximize the likelihood of generating our model.
- We are trying to figure out what model best describes our data. Having domain knowledge can help, in these circustances.

Desireable Effects: (if you have a large sample size)

- · As the sample size increases, we reach an unbiased variance estimator
- The ability to generate a normal distribution and confidence intervals can be used with large enough samples
- The ability to form hypothesis testing

Drawbacks:

- If you have small sample size there will be bias
- · Can be a computational hog

Each probability distribution will have unique parameters. As the parameters change, a different probability distribution is generated.

• The parameters for each model will fall into a family of distributions based on the parameters you have. The parameters here will be (μ, σ)

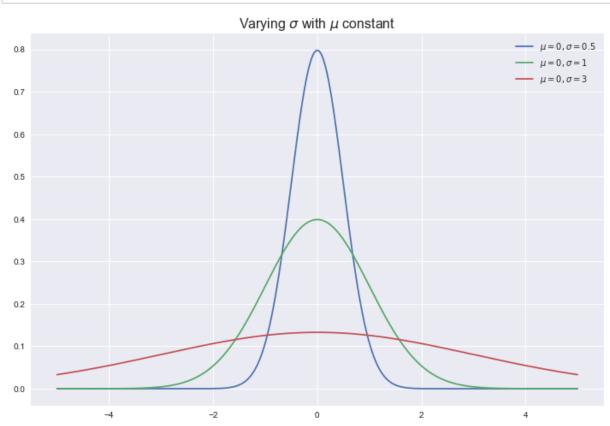
For example you may have a normal distribution, but the mean (μ) and standard deviation (σ) are not the same from one model to the other.

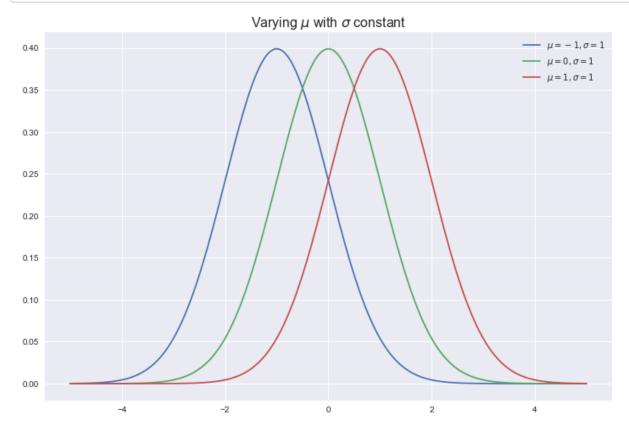
• Therefore, you may have a family of normal distributions with varying parameters.

Plot showing different normals and explain as family of functions

• These two plots reflect the parameters of a Normal Distribution you may want to estimate. If this were a different problem you would evaluate other parameters and which may not be $[\mu, \sigma]$

```
In [1]:
        import numpy as np
        import matplotlib.pyplot as plt
        %matplotlib inline
        plt.style.use('seaborn') # pretty matplotlib plots
        plt.rcParams['figure.figsize'] = (12, 8)
        x = np.linspace(-5, 5, 5000)
        mu = [0,0,0]
        sigma = [.5, 1, 3]
        y_=[]
        for i in range(len(sigma)):
            y = (1 / (np.sqrt(2 * np.pi * np.power(sigma[i], 2)))) * \
            (np.power(np.e, -(np.power((x - mu[i]), 2) / (2 * np.power(sigma[i],
        2)))))
            y_.append(y)
        labels=[('$\mu=0,\sigma=0.5$'),('$\mu=0,\sigma=1$'),('$\mu=0,\sigma=3$'
        ) ]
        for i in range(len(y_)):
            plt.plot(x,y_[i],label=labels[i])
            plt.title('Varying $\sigma$ with $\mu$ constant', size=16)
            plt.legend();
```





In the real world we are given some data that we observed, and need to find some model of interest that will best describe our data using a PDF that will most likely represent our data.

• Realistically, we do not always have the option of transforming our data (log,reciprocal,square root, etc) to save the day.

Instead we have: $L(\theta; x_i) = f(x; \theta)$

 $L(heta;x_i)$: This means that we have the likelihood of some true parameter theta given the observed data X

well what is the right side then?: it is our PDF and there is a distinction

 $f(x; \theta)$: is a function of data given a particular set of parameters

I am using the semicolon to denote parameters with respect to X instead of confusing conditional probability at this point, which would have the vertical bars

Math of MLE:

Nasty Truth:

• First, there doesn't have to be a unique value of the MLE

Second, there may not be a value that exists for the MLE

Hint: because you need to think about concavity, this is a problem if we are dealing with machine learning algorithms and a bit topic.

Super Fun Happy Times, wait what?

S.2.7

MaTh TiMe Y'aLL:

$$L(\theta; x_i) = f(x; \theta)$$

$$f(x; \theta) = \prod f(X|\theta) =$$

$$f(x_1, x_2, x_3 \dots x_n | \theta) =$$

$$f(x_1|\theta) * f(x_1|\theta) * f(x_2|\theta) * \dots f(x_n|\theta)$$

But, now the trickery $^-\setminus_(^)_/^-$:

We will decide to do the Log because, it will get rid of our multiplications and simply everything. Also, since the log is strickly increasing it shouldn't effect our function.

• We will need to find the Max/Min, hince we will need the first derivative with respect to θ and equate to zero.

$$log(L(\theta; x_i)) = log[f(\theta|x_1) * f(\theta|x_1) * f(\theta|x_2) * \dots f(\theta|(x_n))]$$

$$= log(f(\theta|x_1)) + log(f(\theta|x_1)) + log(f(\theta|x_2)) + \dots log(f(\theta|x_n))$$

$$= \sum_{i=1}^{n} log(f(\theta|x))$$

Derivative Time

First Derivative:
$$\frac{\partial ln(L(\theta|x))}{\partial \theta_i} = 0$$

Second Derivative:
$$\frac{\partial^2 ln(L(\theta|x))}{\partial \theta_i^2} < 0$$

At this point if you have a suspected distribution, you would replace with your distribution function. There are times when:

Real World: Either the data are highly paramertized, non-convex (are a few reasons) and you will need to use optimazation techinues to attempt solving these problems with approximation methods.

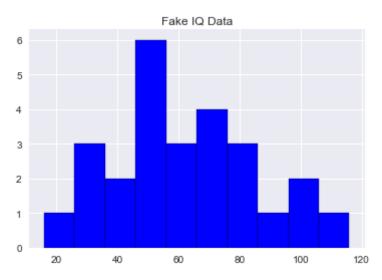
Examples: Newton's method, Fisher scoring, various conjugate gradient-based approaches, steepest descent, Nelder-Mead type (simplex) approaches, BFGS and a wide variety of other techniques (from: https://www.analyticsvidhya.com/blog/2018/07/introductory-guide-maximum-likelihood-estimation-case-study-r/))

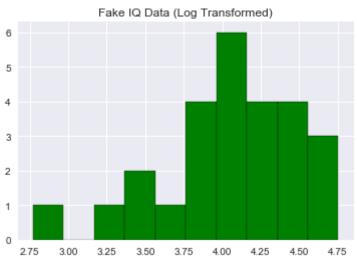
a great resource: https://www.stat.tamu.edu/~suhasini/teaching613/chapter2.pdf

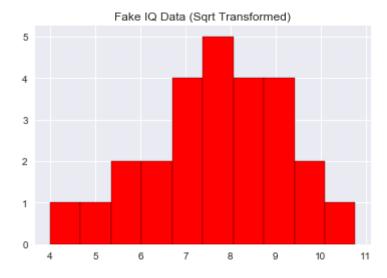
Assume we have some (*fake*) data based on the preceived IQ for people who vote for a particular president

In [122]: # Histogram with density plot import seaborn as sns import matplotlib.mlab as mlab %matplotlib inline fake_data=[30,40,62,61,85,80,16,55,26,53,51,70,116, 66,51,101,99,88,71,55,51,35,62,71,81,44] plt.hist(fake_data, color = 'blue', edgecolor = 'black') plt.title('Fake IQ Data') plt.subplots() plt.hist(np.log(fake_data), color = 'green', edgecolor = 'black') plt.title('Fake IQ Data (Log Transformed)') plt.subplots() plt.hist(np.sqrt(fake_data), color = 'red', edgecolor = 'black') plt.title('Fake IQ Data (Sqrt Transformed)') # plt.subplot() # plt.hist(np.log(fake_data), color = 'green', edgecolor = 'black') # plt.title('Fake IQ Data (Log Transformed)')

Out[122]: Text(0.5, 1.0, 'Fake IQ Data (Sqrt Transformed)')







Lets look at this a little more clearly:

```
In [296]: fig, axarr = plt.subplots(2, 1, figsize=(12, 8))

sns.distplot(fake_data,ax=axarr[0])
plt.suptitle('Fake IQ Data: [original]',size=14)

sns.distplot(np.sqrt(fake_data),ax=axarr[1])
plt.title('Fake IQ Data: [sqrt transformed]',size=14)

plt.show()
```

Fake IQ Data: [original]



transform data and looks close? What now?

- Well, lets consider that maybe we think these data reflect a normal distribution.
 - We could do a comparison after finding the parameters and see how it matches up.
 - We also, may end up in a situation where we need to reconsider our distribution as well. (which is) very likely in the real world.

For simplicity we are assuming Normal distribution:

$$f(x_1, x_2, x_3 \dots x_n; \mu, \sigma) =$$

Replace with distribution:

$$P(X; \mu, \sigma) = \frac{1}{\sigma\sqrt{(2\pi)}} exp(-\frac{(x-\mu)^2}{2\sigma^2}) =$$

$$\prod \frac{1}{\sigma\sqrt{(2\pi)}} exp(-\frac{(x-\mu)^2}{2\sigma^2}) =$$

Expand:

$$\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_1-\mu)^2}{2\sigma^2})*\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_2-\mu)^2}{2\sigma^2})*\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_3-\mu)^2}{2\sigma^2})*...\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_n-\mu)^2}{2\sigma^2})$$

Now take the Log:

$$ln(L(\theta; X)) = ln(\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_1-\mu)^2}{2\sigma^2})) + ln(\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_2-\mu)^2}{2\sigma^2})) + ln(\frac{1}{\sigma\sqrt{(2\pi)}}exp(-\frac{(x_3-\mu)^2}{2\sigma^2})) + \dots ln(\frac{1}{\sigma\sqrt{(2\pi)}exp(-\frac{(x_3-\mu)^2}{2\sigma^2})) + \dots ln(\frac{1}{\sigma\sqrt{(2\pi)}exp(-\frac{(x_3-\mu)^2}{2\sigma^2})) + \dots ln(\frac{1}{\sigma\sqrt{(2\pi)}exp(-\frac{(x_3-\mu)^2}{2\sigma^2})) + \dots ln(\frac{(x_3-\mu)^2}{2\sigma^2}) + \dots ln(\frac{(x_3-\mu)^2}$$

Simplify:

$$ln(L(\theta;X)) = ln(\frac{1}{\sigma\sqrt{(2\pi)}}) + ln(exp(-\frac{(x_1-\mu)^2}{2\sigma^2}))$$
, this is done for each x_i so I will simplify:

$$= \frac{-1}{2} ln(2\sigma\pi) - \frac{(x_1 - \mu)^2}{2\sigma^2}$$

$$= \frac{-1}{2} ln(2\pi) - \frac{1}{2} ln(\sigma) - \frac{(x_1 - \mu)^2}{2\sigma^2}$$
, Now repeat for a X=[x_1,x_2...x_n]

To speed this up and understand that we have: (n) observations (data points)

$$=\frac{-n}{2}\ln(2\pi)-n*\ln(\sigma)-\frac{1}{2\sigma^2}\sum_{i=1}^n(x_i-\mu)^2$$
 This is what we will take the derivative of

Now the Derivatives:

The first derivative will be the slope of the log-likelihood curve:

$$\frac{\partial L}{\partial \mu} = 0 - 0 - \frac{1}{2\sigma^2} \sum_{i=1}^{n} (x_i - \mu)^2 =$$

$$\frac{1}{\sigma^2} \sum_{i=1}^n x_i - \mu =$$

$$\frac{1}{\sigma^2} \sum_{i=1}^n x_i - n\mu$$

$$\hat{\mu} = \frac{\sum_{i=1}^{n} x_i}{n}$$
 =sample mean

-----Now Standard Deviation-----

short cut:

$$= \frac{-n}{2}ln(2\pi) - n * ln(\sigma) - \frac{1}{2\sigma^2} \sum_{i=1}^{n} (x_i - \mu)^2$$

$$\frac{\partial L}{\partial \sigma} = 0 - \frac{n}{\sigma} + \frac{1}{\sigma^3} \sum_{i=1}^n (x_i - \mu)^2$$
 =, set equal to zero

$$-\frac{n}{\sigma} + \frac{1}{\sigma^3} \sum_{i=1}^{n} (x_i - \mu)^2 = 0$$

Manipulate:

$$\frac{1}{\sigma^3} \sum_{i=1}^n (x_i - \mu)^2 = \frac{n}{\sigma}$$

$$\hat{\sigma} = \sqrt{rac{\sum_{i=1}^{n}(x_i - \mu)^2 = rac{n}{\sigma}}{n}}$$
, biased estimator

side note: the (n-1) will be unbiased look into proof, second webpage below

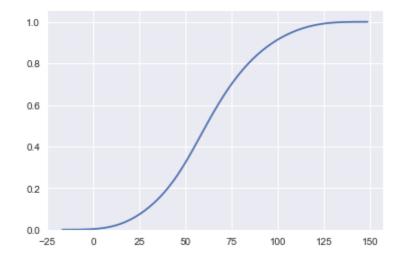
https://machinelearningmastery.com/probability-density-estimation/ (https://machinelearningmastery.com/probability-density-estimation/)

https://daijiang.name/en/2014/10/08/mle-normal-distribution/ (https://daijiang.name/en/2014/10/08/mle-normal-distribution/)

Check the CDF: so we can get an idea of if these data will fit a Normal Distr

```
In [511]: kwargs = {'cumulative': True}
# sns.distplot(fake_data, hist_kws=kwargs, kde_kws=kwargs)
sns.kdeplot(fake_data,cumulative=True)
```

Out[511]: <matplotlib.axes._subplots.AxesSubplot at 0x1a424b5208>

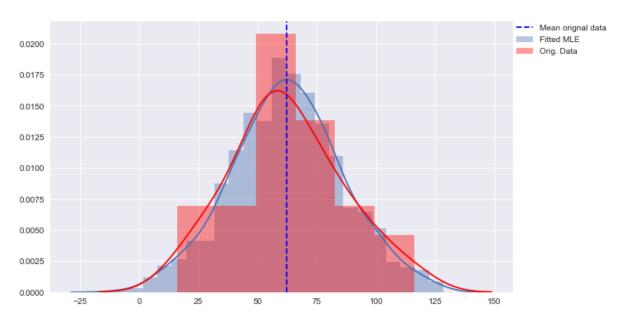


We can directly compute the μ & σ from the data, but this doesn't mean that we are actually representing the population the data came from.

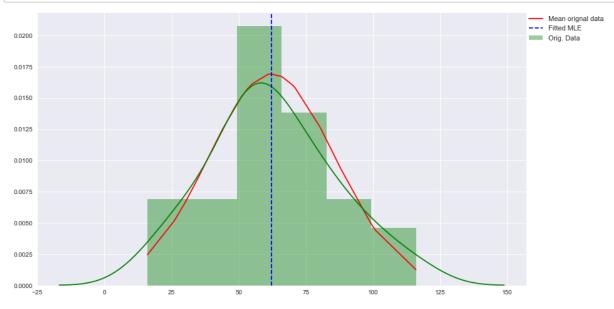
• Hence, the use of MLE.

```
In [452]: print(np.mean(fake_data),np.std(fake_data))
          fig dims = (10, 6)
          fig, ax = plt.subplots(figsize=fig_dims)
          sns.distplot(np.random.normal(np.mean(fake_data),np.std(fake_data),1000
          ))
          # plt.legend('MLE')
          sns.distplot(fake_data,color='red')
          plt.axvline(np.mean(fake data),color='b', linestyle='--')
          plt.legend(['Mean orignal data','Fitted MLE','Orig. Data'],bbox to ancho
          r=(1, 1),
                      loc=2, borderaxespad=0.) #,fontsize=25
          plt.show()
          # sigma=np.std(fake data)
          # mu=np.mean(fake data)
          # vals=[]
          # for x in sorted(fake data):
                  norm d=1/(sigma*np.sqrt(2*np.pi))*np.exp(-(x-mu)^2/2*sigma^2)
                norm d=1/(sigma*np.sqrt(2*np.pi))*np.exp(-np.power(x - mu, 2.) / mu)
           (2 * np.power(sigma, 2.)))
                vals.append(norm d)
          # sorted(fake data)
```

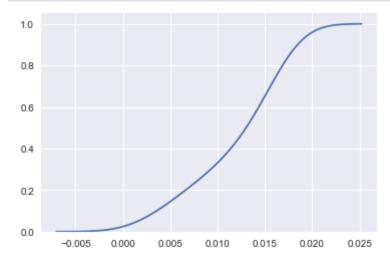
62.30769230769231 23.57601328398981



```
In [508]:
          import numpy as np
          from scipy.stats import norm
          import matplotlib.pyplot as plt
          import scipy.stats as stats
          fig dims = (14, 8)
          fig, ax = plt.subplots(figsize=fig_dims)
          meany=np.mean(fake_data)
          sd_=np.std(fake_data)
          pdf = stats.norm.pdf(sorted(fake_data), meany, sd_)
          plt.plot(sorted(fake_data), pdf,color='red') # including h here is cruci
          al
          sns.distplot(fake_data,color='green')
          plt.axvline(np.mean(fake_data),color='b', linestyle='--')
          plt.legend(['Mean orignal data','Fitted MLE','Orig. Data'],bbox to ancho
          r=(1, 1),
                     loc=2, borderaxespad=0.,fontsize=11)
          plt.show()
```



```
In [513]: sns.kdeplot(pdf,cumulative=True)
   plt.show()
```



Last Thoughts:

- You can't always, assume your data will fit an exact model
- · Approximation methods are usually a good call on real data
- Do not blindly, believe everything you hear!

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