## Likelihood

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January 18, 2021

Biol 520C: Statistical modelling for biological data

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- Practical 02 is now unlocked on canvas, and due next Tuesday.

## Review

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$$Pr\{A|B\} = Pr\{A,B\}/Pr\{B\}$$

**A** — The conditional probability of A given B occurred.

**B** — The conditional probability of B given A occurred.

**C** — Bayes' theorem.

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And I told you that this was important for fitting a model to data.

## Likelihood



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For example, we can extend this to indicate that the probability of observing data  $Y_i$  given parameter value p is  $Pr\{Y_i|p\}$ .



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The subscript i indicates that there are multiple possible outcomes, but only one parameter value p.

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In other words, we have a mathematical description for saying "What is the probability of observing our data  $(Y_i)$  given our hypothesis (rate parameter  $= \lambda$ )?"

### Clicker question: ii



Let's say I'm conducting a bird survey in my back yard. What's the probability of counting 2 crows given a rate parameter of 6?

NB: The Poisson distribution has a probability mass function of  $\frac{\lambda^k e^{-\lambda}}{k!}$ 

$$\mathbf{A} - \approx 0.27$$

$$C -\approx 0.16$$

$$\mathbf{B} - \approx 0.012$$

$$D - \approx 0.045$$

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This is where the concept of likelihoods comes in, and we write this:

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Note how here the data are not subscripted (we only observed one outcome), but there are multiple possible parameter values  $p_m$ .





The key distinction between likelihoods and probabilities is that with probabilities the hypothesis is *known*, but the data are *unknown* whereas with likelihoods the data are *known* but the hypothesis is *unknown*.



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$$\mathcal{L}(p_m|Y) = c \operatorname{Pr}\{Y|p_m\}$$



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Because we're typically interested in relative likelihoods (not exact values), the proportionality constant, c, can be set to 1



Ok, cool, but so what?





If 
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For example, if we come back to those 2 crows I counted earlier we can now ask (and answer!) what's the more likely  $\lambda$ , 4, or 6?



If  $\mathcal{L}(p_m|Y) = \Pr\{Y|p_m\}$ , and we have some data, and if we make some distributional assumptions, we have a way of calculating the likelihood of specific parameter values, and we can generalise this to any number of parameters and any distribution!

For example, if we come back to those 2 crows I counted earlier we can now ask (and answer!) what's the more likely  $\lambda$ , 4, or 6?

$$\mathcal{L}(\lambda=4|2) = \frac{4^2e^{-4}}{2!} \approx 0.15$$
  $\mathcal{L}(\lambda=6|2) = \frac{6^2e^{-6}}{2!} \approx 0.045$ 

Remember that the Poisson distribution has a PMF given by  $\frac{\lambda^{k}e^{-\lambda}}{k!}$ 



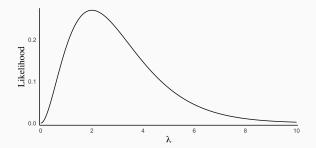


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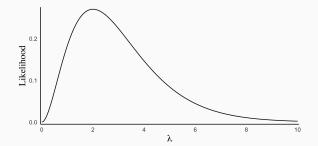
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For our 2 crows, a plot of the likelihood as a function of  $\lambda$  looks like this:



The value of  $\lambda$  that maximises the likelihood is the maximum likelihood estimate (MLE) of our parameter value





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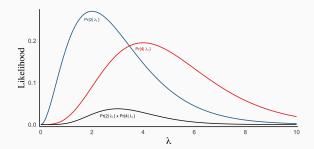
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If we assume our observations are independent, then  $\Pr\{A, B|p_m\} = \Pr\{A|p_m\} \times \Pr\{B|p_m\}$ 



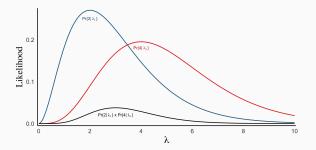


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Here,  $\hat{\lambda}=3$  is the maximum likelihood estimate (MLE)





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$$\mathcal{L}(\lambda|x_i) = \prod_{i=1}^n \frac{\lambda^{x_i} e^{-\lambda}}{x_i!}$$





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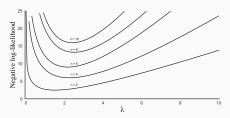
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In most cases, as the sample size increases, the negative log-likelihood function becomes increasingly peaked around its maximum



#### MLE in action



Let's say I go out and measured the height of ten people. I come back with  $\{171,168,180,190,169,172,162,181,177,181\}$  (in cm).

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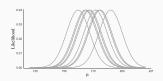
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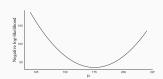
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Plugging in data the resulting likelihood functions look like this:







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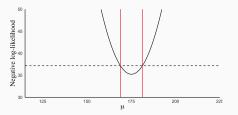
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In the likelihood framework, we can use the likelihood profile to identify the 95% confidence intervals on our estimated parameter. E.g., a simple rule is to place the bounds within 1.92 of the minimum log-likelihood.



So we can say that  $\mu=175.1$  with 95% CIs of  $\sim169-181$ 

#### MLE and Cls.

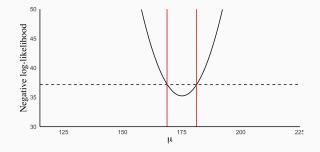


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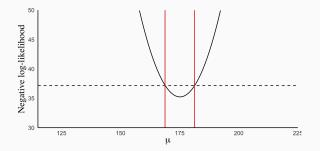


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More data narrows the CIs (and vice versa, less data increases the amount of uncertainty in the MLE)

## MLE and fitting a line to data



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We started our detour into probability theory and maximum likelihood because we learned that the least-squares approach didn't provide any way of understanding how stochasticity entered into a system.

We've now picked up enough of the basics to go back to our linear regression problem and fit a line to some data as probalists.



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and  $\varepsilon_i$  is our Gaussian distributed error with a mean of 0 and variance of  $\sigma^2$ , whose PDF is given by:

$$\frac{1}{\sigma\sqrt{2\pi}}e^{-\frac{1}{2}\left(\frac{x-\mu}{\sigma}\right)^2}$$



Given our dataset  $(x_1, y_1), (x_2, y_2), ..., (x_n, y_n)$ 



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For convenience, we work with negative log-likelihoods, which changes this to:

$$\mathcal{L}(x_i; \beta_0, \beta_1, \sigma^2 | y_i) = n(\log(\sigma) + \frac{1}{2}\log(2\pi)) + \frac{1}{2\sigma^2} \sum_{i=1}^{n} (y_i - (\beta_0 + \beta_1 x_i))^2$$





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After some math that we won't go over, we get the following 3 estimators:

$$\hat{\beta}_1 = \frac{\sum\limits_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})}{\sum\limits_{i=1}^{n} (x_i - \bar{x})^2}$$

$$\hat{\beta}_0 = \bar{y} - \hat{\beta}_1 \bar{x}$$

$$\hat{\sigma}^2 = \frac{1}{n} \sum_{i=1}^{n} (y_i - (\hat{\beta}_0 + \hat{\beta}_1 x_i))^2$$





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i) A description of the system's stochastic component



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So what did we gain by making a Gaussian-noise assumption and estimating the parameters via maximum likelihood?

- i) A description of the system's stochastic component (which allows us to make more realistic predictions)
- ii) A framework for placing confidence intervals on our parameter estimates using the likelihood profiles (which allows us to make formal statistical inference on the parameters)

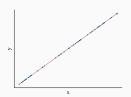
# Stochasticity and predictions



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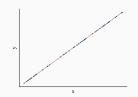


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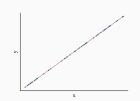


But real systems are full of stochasticity, so the predictions of purely deterministic models are almost certainly going to be wrong.

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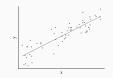


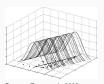
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But real systems are full of stochasticity, so the predictions of purely deterministic models are almost certainly going to be wrong.

With a stochastic component, outcomes are variable and models provide a distribution of the values that  $y_i$  can be expected to take





Source: Zuur et al. 2009





For example, let's say we know that the wingspan (in mm) a species of insect is proportional to its mass (in g), with a slope of 5 and intercept of 10.



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How informative is this? What if we measure an insect with weight = 2 and wingspan = 21? What does this tell us about our model?

# Stochasticity and predic. examp. cont.



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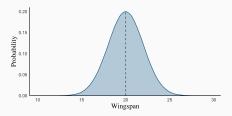
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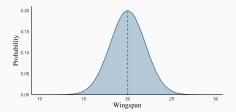


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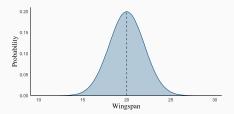
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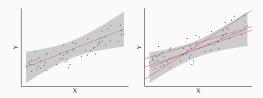
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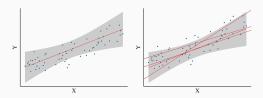




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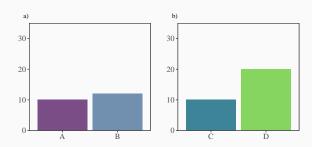
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What happens to prediction intervals when  $n \to \infty$ ? Which will be wider, confidence intervals, or prediction intervals?

## Clicker question: iii

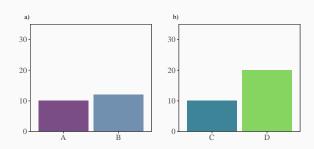




$${\bf C}$$
 — Both a and b

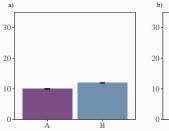
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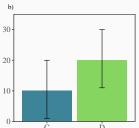




## Clicker question: iv



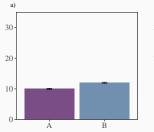


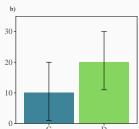


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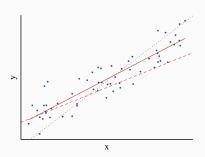
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Which of these models are different from one another?

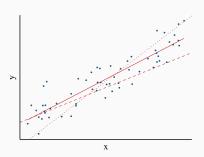
A — The solid from the dashed

B — the dotted from the solid

**C** — The solid from the dotted, and the dotted from the dashed, but not the solid from the dashed.

**D** — I DON'T KNOW.

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The answer depends on the amount of information in our data, the (un)certainty in our estimate, and the shape of our likelihood profile.

