Bayesian statistics with R

2. The likelihood

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Likelihood

Context

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- In the real world, it is usually the other way around.

A question of interest might be for example:

We have observed 3 births by a female during her 10 breeding attempts. What does this tell us about the true probability of getting a successful breeding attempt from this female? For the population?

- We don't know what the probability of a birth is.
- But we can calculate the probability of getting our data for different values:

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#> [1] 0.05739563
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- We don't know what the probability of a birth is.
- But we can see what the probability of getting our data would be for different values:

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dbinom(x=3,size=10,prob=0.9)
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- We don't know what the probability of a birth is.
- But we can see what the probability of getting our data would be for different values:

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dbinom(x=3,size=10,prob=0.25)
#> [1] 0.2502823
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dbinom(x=3,size=10,prob=0.1)
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dbinom(x=3,size=10,prob=0.9)
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dbinom(x=3,size=10,prob=0.25)
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So we would be more likely to observe 3 births if the probability is 0.25 than 0.1 or 0.9.

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- This reasoning is so common in statistics that it has a special name:
- **The likelihood** is the probability of observing the data under a certain model.
- The data are known, we usually consider the likelihood as a function of the model parameters $\theta_1, \theta_2, \dots, \theta_p$

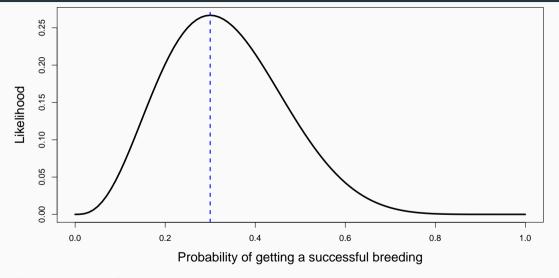
$$L = P(\theta_1, \theta_2, \dots, \theta_p \mid \mathsf{data})$$

Likelihood functions

We may create a function to calculate a likelihood:

```
lik.fun <- function(parameter){</pre>
  11 <- dbinom(x=3, size=10, prob=parameter)</pre>
  return(11)
lik.fun(0.3)
#> [1] 0.2668279
lik.fun(0.6)
#> [1] 0.04246733
```

Maximize the likelihood (3 successes ot of 10 attempts)



The maximum of the likelihood is at value 0.3

Maximum likelihood estimation

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- There is always a set of parameters that gives you the highest likelihood of observing the data, and this is the MLE.
- These can be calculated using:
 - Trial and error (not efficient!).
 - Compute the maximum of a function by hand (rarely doable in practice).
 - An iterative optimization algorithm: ?optim in R.

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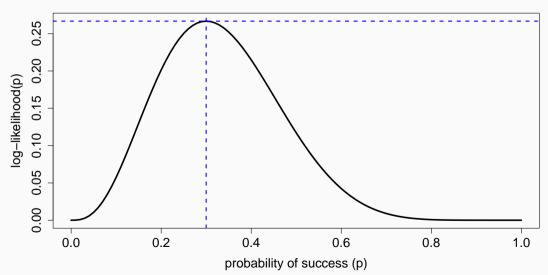
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- Here, the MLE is the proportion of observed successes.

```
lik.fun <- function(parameter) dbinom(x=3, size=10, prob=parameter)
# ?optimize
optimize(lik.fun,c(0,1),maximum=TRUE)
#> $maximum
#> [1] 0.3000157
#>
#> $objective
#> [1] 0.2668279
```

Use optim when the number of parameters is > 1.

Binomial likelihood with 3 successes ot of 10 attempts

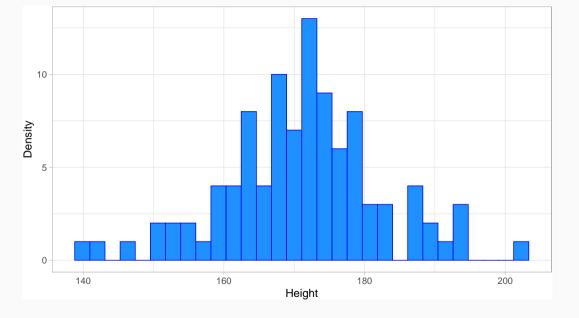


Your turn

MLE of the parameters of a Normal distribution

• Assume we have collected data on the height of 100 people:

```
# set seed for random numbers
set.seed(2020)
# simulate data from Normal distribution
n <- 100
height <- rnorm(n, mean=170, sd=10)</pre>
```



- We consider a Normal distribution for the model.
- Compute the MLE of the parameters of the Normal distribution.
- Hint: Use functions optim() and dnorm()

Solution

R code

• Write a function for the likelihood of a Normal distribution with parameters mean μ and standard deviation σ :

```
negloglik <- function(theta, data) {
  mu <- theta[1]
  sigma <- theta[2]
  x <- data
  -sum(dnorm(x, mean = mu, sd = sigma, log = TRUE))
}
negloglik(theta = c(150,1), height)
#> [1] 28530.45
```

Minimiiiiize

```
fit <- optim(par = c(1,1), fn = negloglik, data = height)
fit
#> $par
#> [1] 171.05358 11.16656
#>
#> $value
#> [1] 382.9207
#>
#> $counts
#> function gradient
#>
        135
                  NA
#>
#> $convergence
#> \[ 17 \ 0 \]
```

