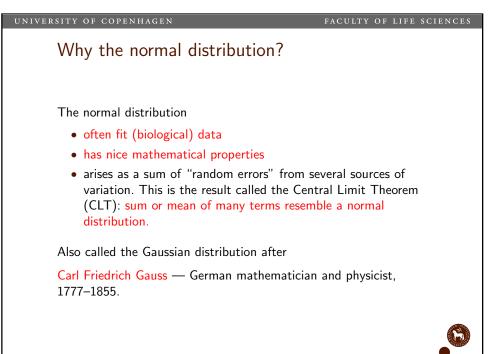


Program

- The normal distribution (continued)
- Density function
- Comparing data with the normal distribution
- Scaling and standardization
- The standard normal distribution
- The normal distribution in R
- Sums of normals are again normal
- Normal probability calculations
- Probabilities of centered intervals
- Summary of main points

lide 2 — Statistics for Life Science (Week 3-1 2010) — The Normal distribution





Weights of crabs

• Weights of 162 crabs of a

certain age: y₁,..., y₁₆₂. • R: $\bar{y} = 12.76$, s = 2.25

• Histogram normalized to have total area 1

• Curve for *f*, where *f* is the

density of the normal

distribution

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Density and probabilities

Density for the normal distribution with mean μ and standard deviation $\sigma:$

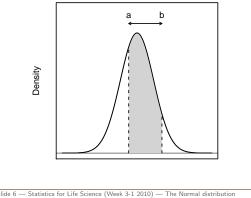
$$f(y) = \frac{1}{\sqrt{2\pi\sigma^2}} \exp\left(-\frac{1}{2\sigma^2}(y-\mu)^2\right)$$

• Scaling preserves normality: If Y is normal then $a + b \cdot Y$ is

normal with the "natural" mean $a + b\mu$ and standard

• Standardization: In particular, the standardized version,

 $Z = \frac{Y - \mu}{\sigma}$ has mean zero and standard deviation one.

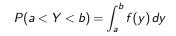


Scaling and standardization

deviation $|b|\sigma$.

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The probability (= population frequency) of the interval between a and b is equal to the area under the density curve within that interval:



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Are the data normally distributed?

Weight of 162 crabs: y_1, \ldots, y_{162} . Sample mean is $\bar{y} = 12.76$ and standard deviation is s = 2.25.

 $f(y) = \frac{1}{\sqrt{2\pi \cdot 2.25^2}} \exp\left(-\frac{1}{2 \cdot 2.25^2} (y - 12.76)^2\right)$

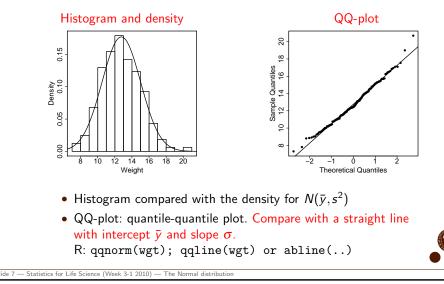
0.15

Density 0.10

0.05

0.00

12 14 16 Weight



The standard normal distribution

Probabilities from $N(\mu, \sigma^2)$ may be converted to probabilities from N(0,1). For example,

$$P(Y \le a) = P\left(\frac{Y - \mu}{\sigma} \le \frac{a - \mu}{\sigma}\right) = P\left(Z \le \frac{a - \mu}{\sigma}\right) = \Phi\left(\frac{a - \mu}{\sigma}\right)$$

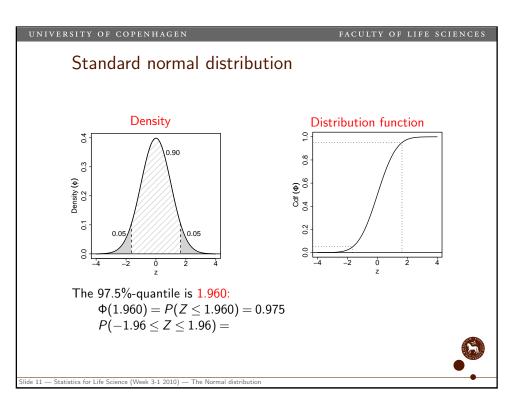
Density of N(0,1):

$$\phi(z) = \frac{1}{\sqrt{2\pi}} \exp\left(-\frac{1}{2}z^2\right)$$

Distribution function — "area to the left of z":

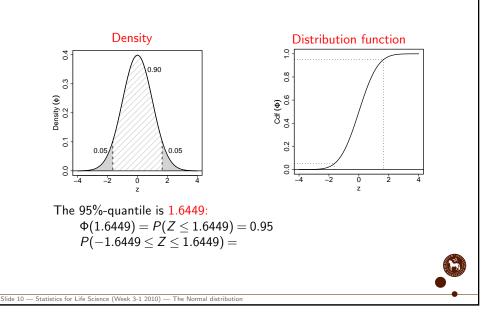
$$\Phi(z) = P(Z \le z) = \int_{-\infty}^{z} \phi(x) \, dx$$

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Standard normal distribution



DIVERSITY OF COPENHAGEN Normal distribution function in R The distribution function for $N(\mu, \sigma)$ is $\Phi_{\mu,\sigma}(x) = P(Y \le x)$, which in R is the function pnorm(..): > pnorm(x, mean= mu, sd=sigma) Examples (crab weights): > pnorm(14, mean= 12.76, sd=2.25) [1] 0.7092212 # Prob. of a value less or equal to 14 > pnorm(14, mean= 12.76, sd=2.25, lower.tail=FALSE) [1] 0.2917888 # Prob. of a value greater than 14



Quantiles in $N(\mu, \sigma)$ are values of the inverse distribution function. In R it is written

> qnorm(q, mean= mu, sd=sigma)

Example (crab weights):

0.75 = Prob. of a value less or equal to ?
> qnorm(0.75, mean= 12.76, sd=2.25)
[1] 14.27760
Answer: 75% of the population has values below 14.28



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Sum of normally distributed variables

Sum of normals is again normal:

If Y_1 and Y_2 are independent and both normal then their sum is again normal with

- mean of sum = sum of means,
- variance of sum = sum of variances,
- BUT NOT standard deviation of sum = sum of standard deviations,

This holds also for the sum of more than two variables.



R: The standard normal distribution

In pnorm(..) and qnorm(..) in R, mu=0 and sigma=1 are default values, corresponding to the standard normal distribution.

- Distribution function: pnorm, fx.
 - > pnorm(1.6449) [1] 0.9500048
- Quantiles: qnorm, fx.
 - > qnorm(0.975) [1] 1.959964

Compare with the table in Appendix C2, p. 400 in the book.

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N-probabilities

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Crab data: assume that crab weight $\sim N(12.76, 2.25^2)$

What is the probability that

- a randomly selected crab weighs at most 14 gram?
- en a randomly selected crab weighs at least 10 gram?
- a randomly selected crab weighs between 10 and 14 gram?
- the sum of the weights of two randomly selected crabs is at least 26 gram?
- > pnorm(1.227)
 [1] 0.8900887
 > pnorm(-1.227)
 [1] 0.1099113
 > pnorm(0.151)
 [1] 0.5600121

> pnorm(0.5511)
[1] 0.7092174
> pnorm(-0.5511)
[1] 0.2907826

