

Tópicos I – Morfometria Geométrica

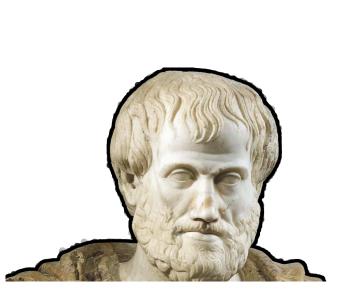


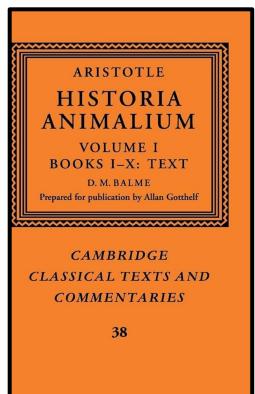


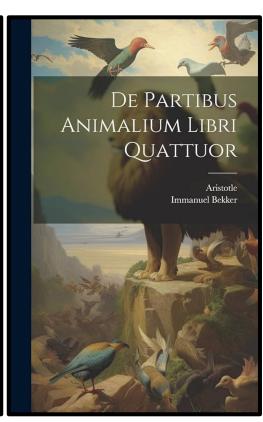


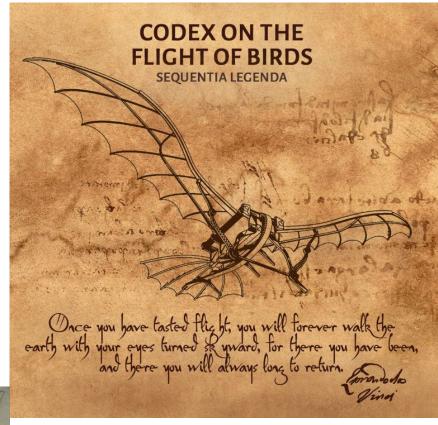


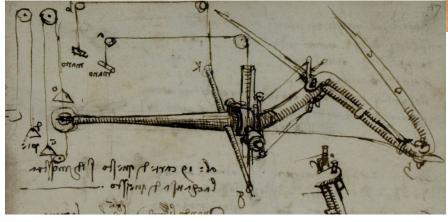
Antiguidade: Aristóteles



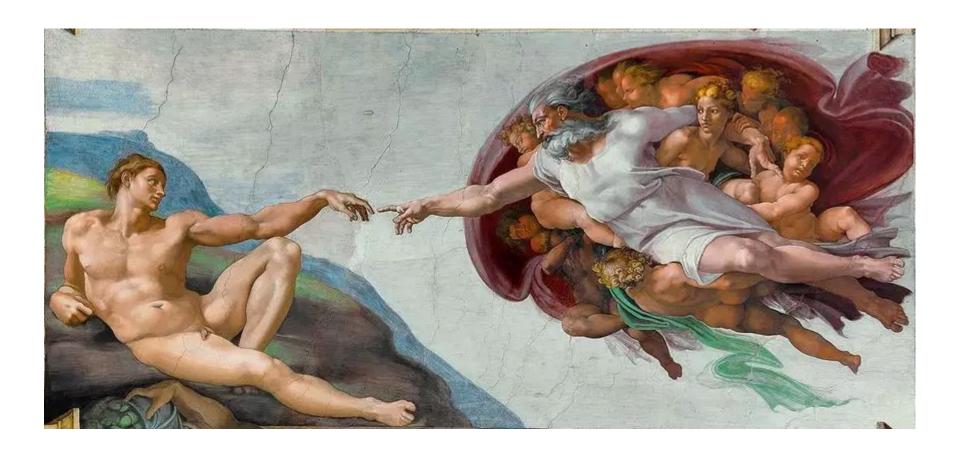


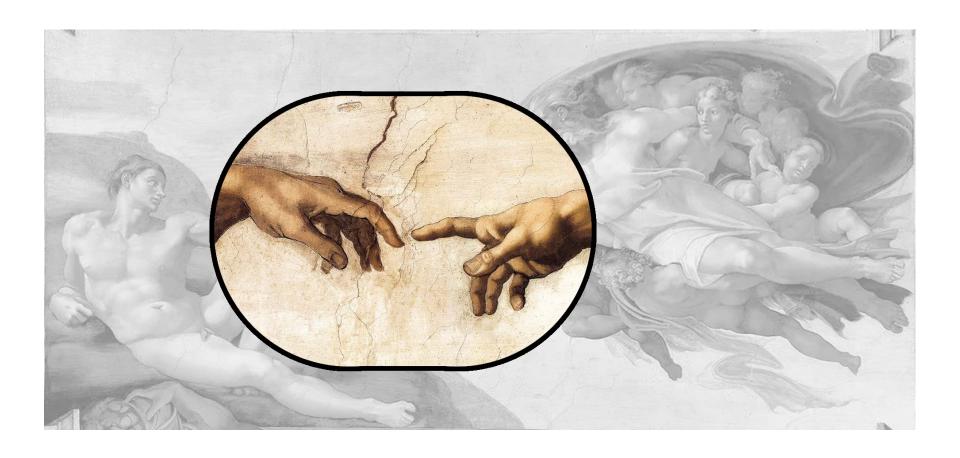


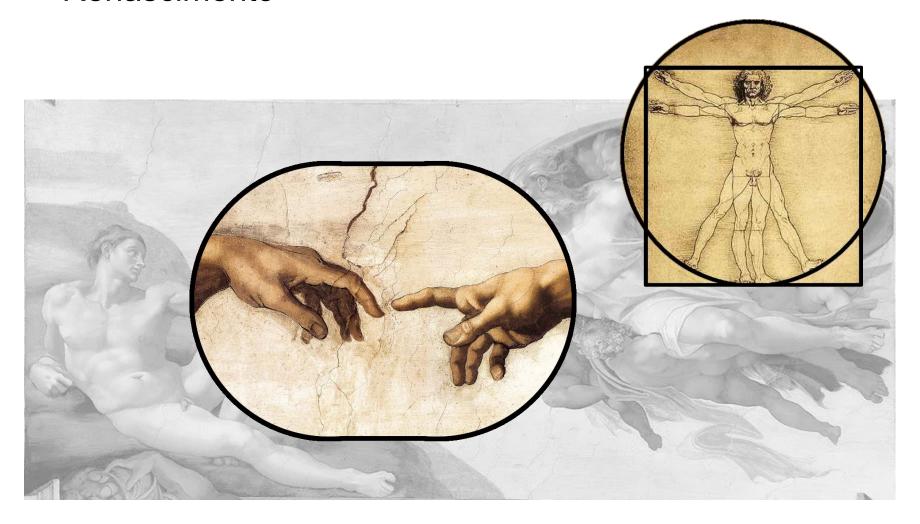




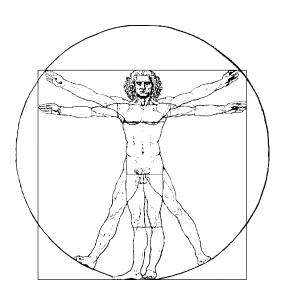


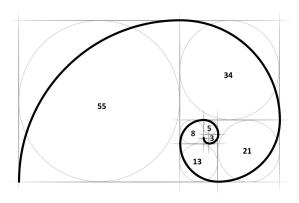




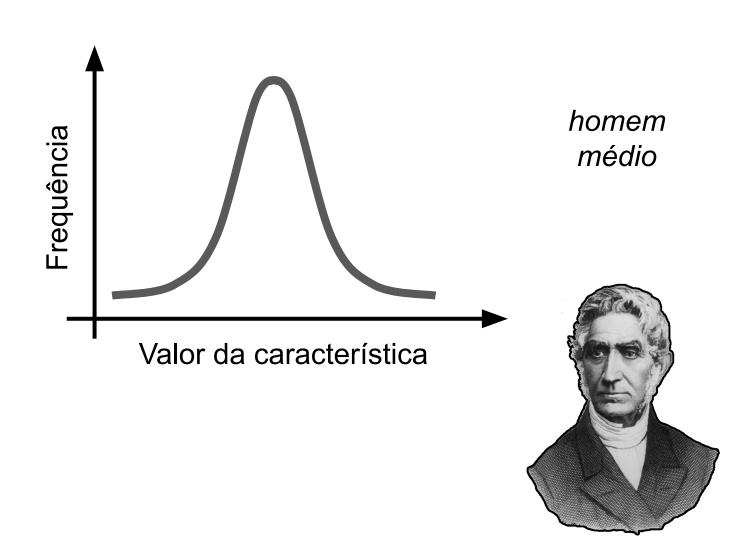


Busca pela compreensão de proporções

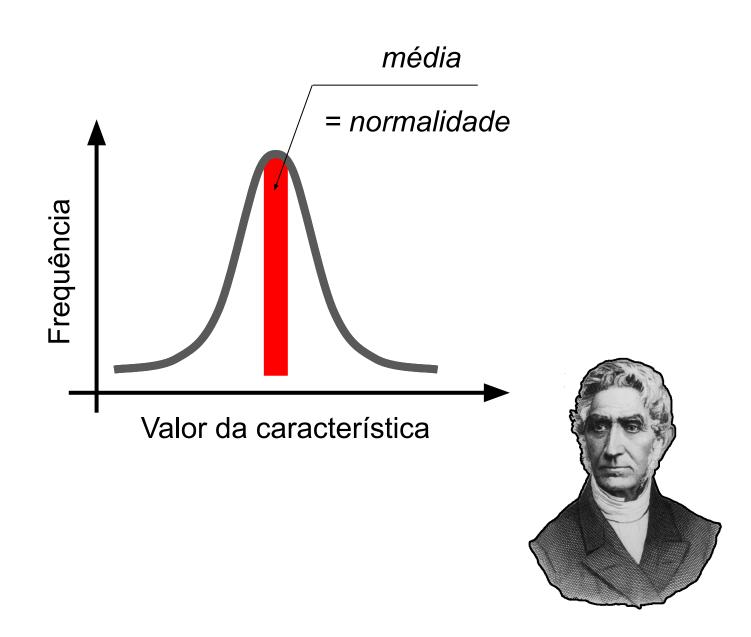




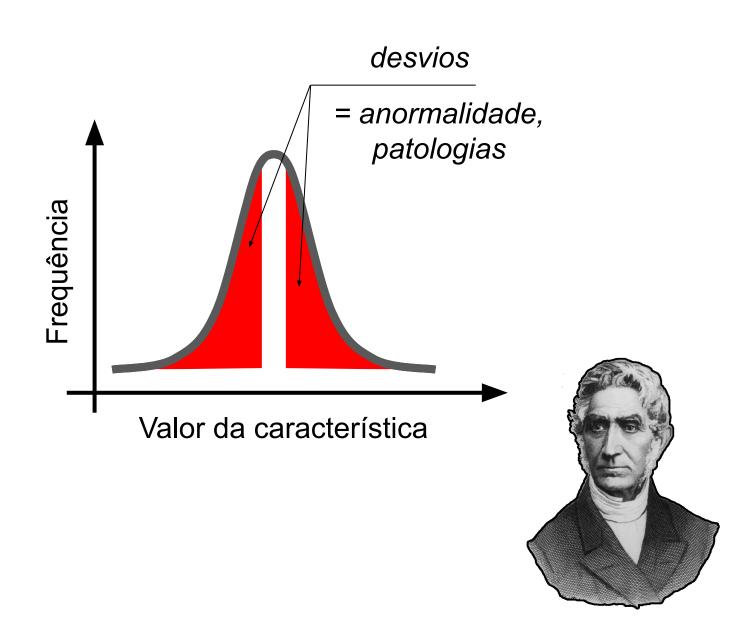
• Século XVII: Adolphe Quételet



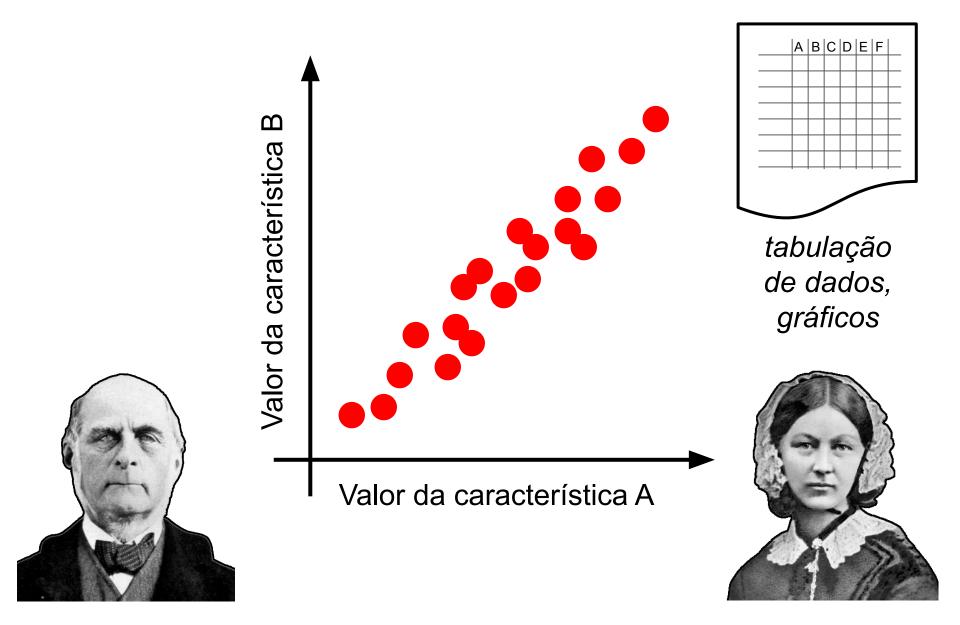
Século XVII: Adolphe Quételet



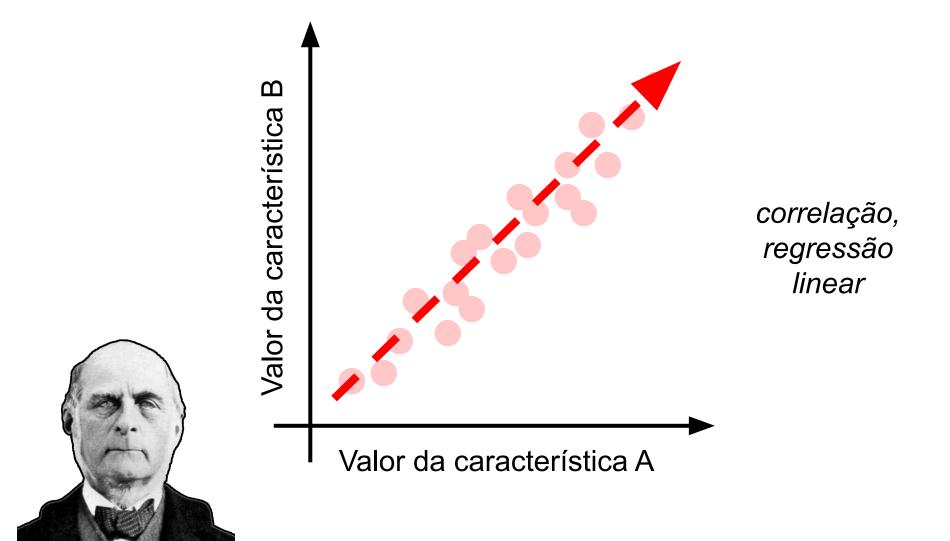
• Século XVII: Adolphe Quételet



Século XIX: Francis Galton, Florence Nightingale



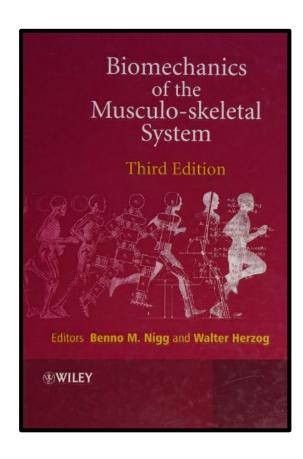
Século XIX: Francis Galton

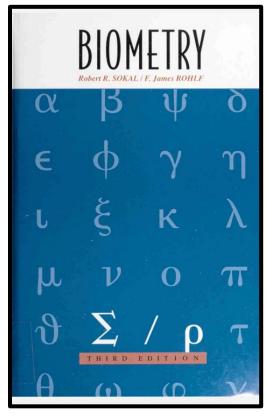


Século XX: Karl Pearson, Ronald A. Fisher







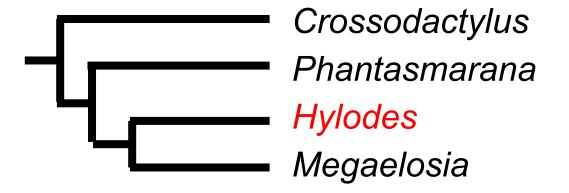


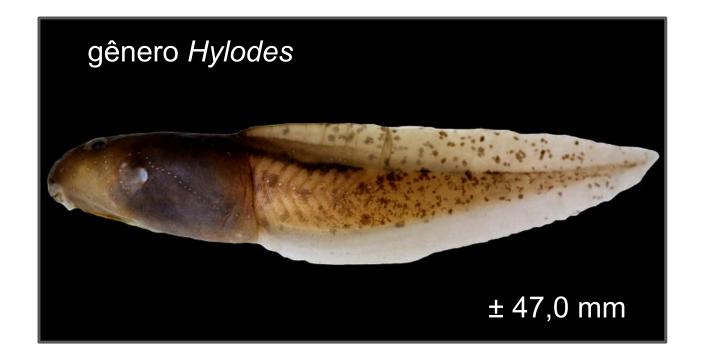
Imagine que tenho dois grupos taxonômicos que:

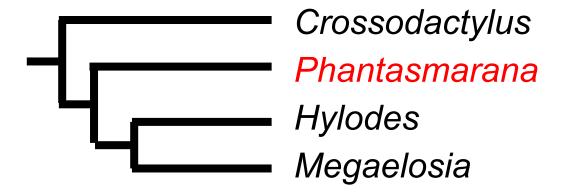
 são próximos filogeneticamente

Imagine que tenho dois grupos taxonômicos que:

 são próximos filogeneticamente muito diferentes no tamanho









Será que essas espécies são morfologicamente diferentes, ou sua diferença é apenas uma questão de escala?

Hora de praticar:



https://imagej.net/software/fiji/downloads

https://cran.r-project.org/

https://posit.co/download/rstudio-desktop/



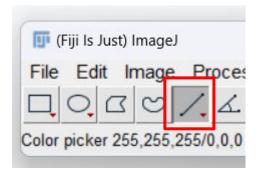
Planilha: shorturl.at/IIfCw

Fotos: shorturl.at/eKIZh



0. File > Open > Navegar até a pasta

1. Clique na Iinha reta



Indique a escala da foto

1. Clique na 2. Analyze > Set Scale

Distance in pixels: 203.5
Known distance: 0.00
Pixel aspect ratio: 1.0
Unit of length: pixel
Click to Remove Scale
☐ Global
Scale: <no scale=""></no>
OK Cancel Help

3. Criar nova reta para a medida desejada > Ctrl + M



Duas abordagens:

1. Usar os resíduos do modelo

```
modelo <- gls(medidas~size)
prcomp(modelo$residuals)</pre>
```

2. Usar uma transformação por média geométrica