

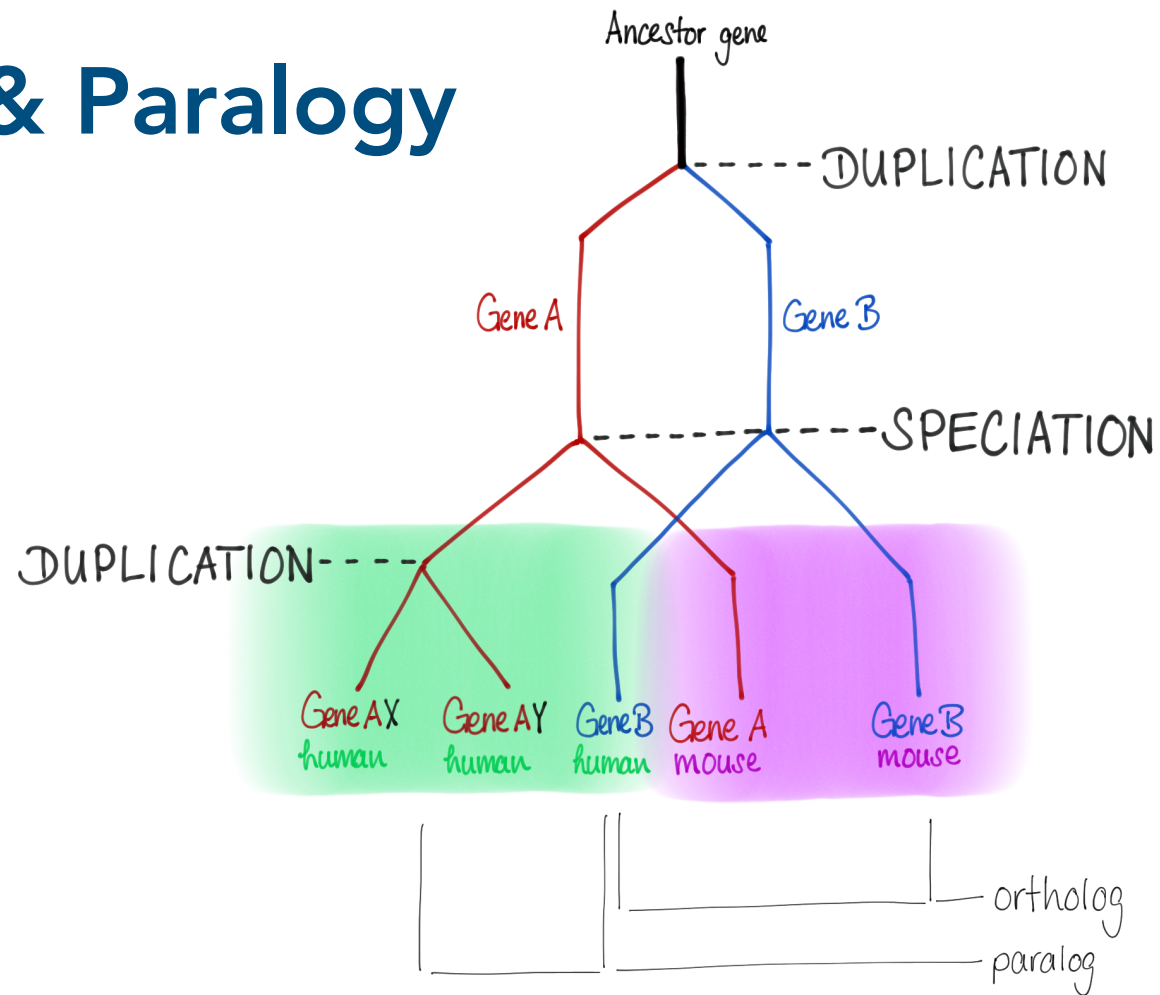
# OrthoInspector: comprehensive orthology analysis and visual exploration

Benjamin Linard, Julie D. Thompson, Olivier Poch, Odile Lecompte

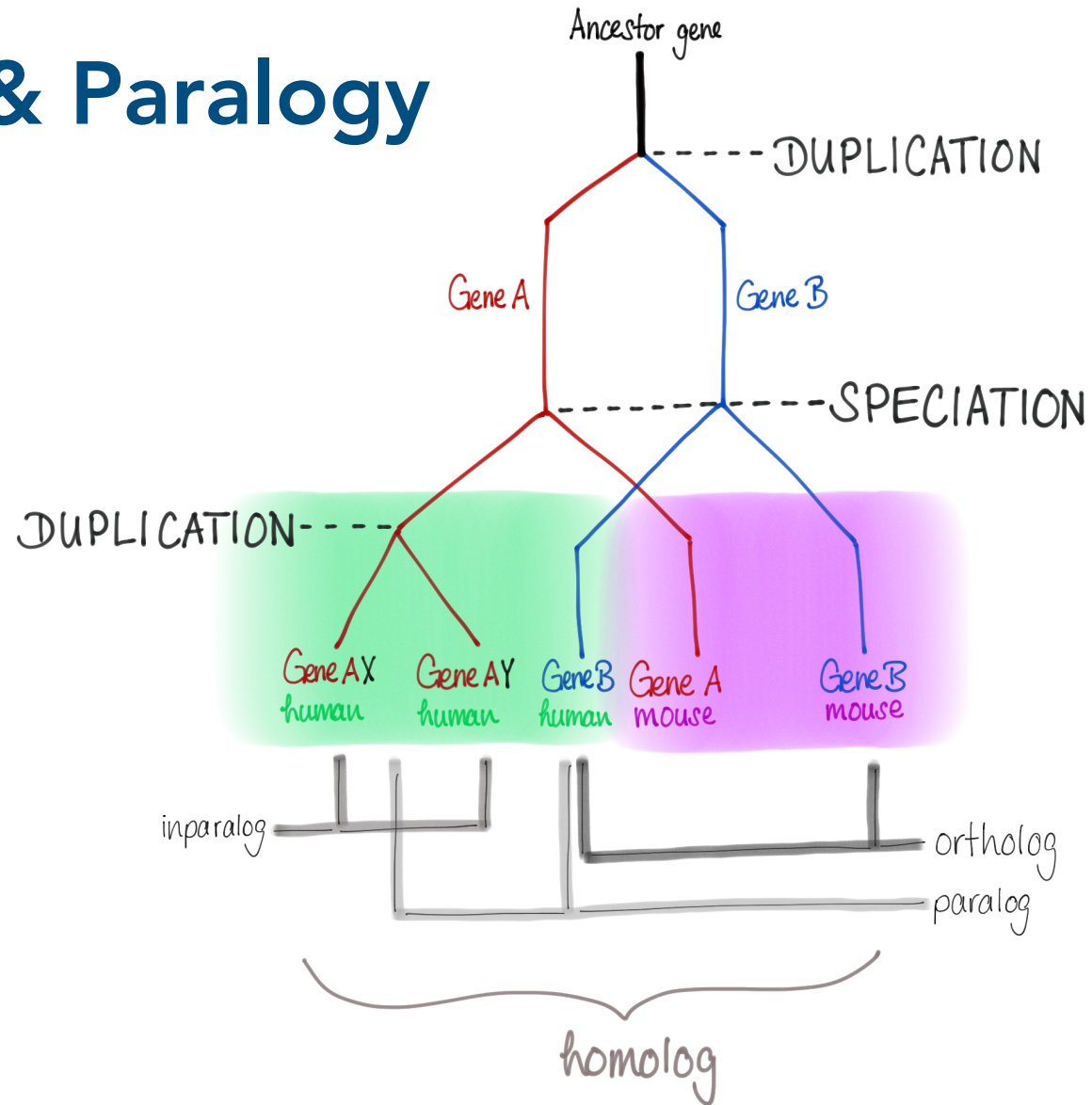
Kassoum Wonogo and Carina Zschammer

# Biological Motivation

# Ortho- & Paralogy



# Ortho- & Paralogy



# Problem & Solution

## Find all Orthologs

Problems of existing methods:

- Primary search for reciprocal best hits in sequence data
- Risk of false negative hits

Solution by OrthoInspector:

- First find inparalog genes and form corresponding groups
- In a second step pairwise comparisons between inparalog groups to find potential (co)-orthologs

# Practical Motivation

# Problem & Solution

## Software Application

Problems of existing methods:

- Complex use: programming knowledge is often required

Solution by OrthoInspector:

- Application via two user interfaces: command line client & graphical interface

# Conclusion

- OrthoInspector is a functional software tool for orthology/inparalogy analyses
- Reliable and safe predictions
  - Well balanced sensitivity and specificity
  - Advantages of different prediction methods should be combined to further improve both sensitivity and specificity
- Very user-friendly



# Thank you for your attention!

Linard, B., Thompson, J.D., Poch, O. *et al.* OrthoInspector: comprehensive orthology analysis and visual exploration. *BMC Bioinformatics* **12**, 11 (2011). <https://doi.org/10.1186/1471-2105-12-11>