## Orthology Analysis part of "Graphen und Netzwerke in der Biologie"

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Sonja Prohaska Orthology Analysis

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#### "A is homologous to B" means...

- A and B are characters (nucleotide sequences, organs,...) of individual organisms I<sub>A</sub> and I<sub>B</sub>, respectively
- A and B are derived from a (last) common ancestor lca(A, B) by descent
- if *A* is a homolog of *B*, *B* is a homolog of *A* (symmetric relation)
- "homologous" in respect to syntax (structure) or sematics (function)?
- o does NOT just mean "A is similar to B"
- similarity might hint at homology
- similarity without common ancestry is called analogy

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# Orthology – Paralogy

- if *A* and *B* drived from the *lca*(*A*, *B*) by **duplication**, *A* and *B* are **paralogous**
- if A and B are homologs and I<sub>A</sub> = I<sub>B</sub> than they are in-paralogs
- if A is paralogous to B and I<sub>A</sub> ≠ I<sub>B</sub> than they are out-paralogs
- if A and B drived from the *lca*(A, B) by **speciation**, A and B are **orthologous**



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# Orthology – Paralogy



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## 1:1, 1:many, many:many orthology



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## Problem of destinguishing orthologs and paralogs

# More information from additional species and about timing of gene duplication and speciation events can change the view.



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