

Swiss Institute of
Bioinformatics

3- Gene and genome duplication

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SIB course Biodiversity bioinformatics 2023



Unil
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Département d'écologie
et évolution

Bgee



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Main types of duplication

Retroposition

- frequent
- most new paralogs dead on arrival

Tandem duplication

- frequent
- can lead to arrays of tandem duplicates

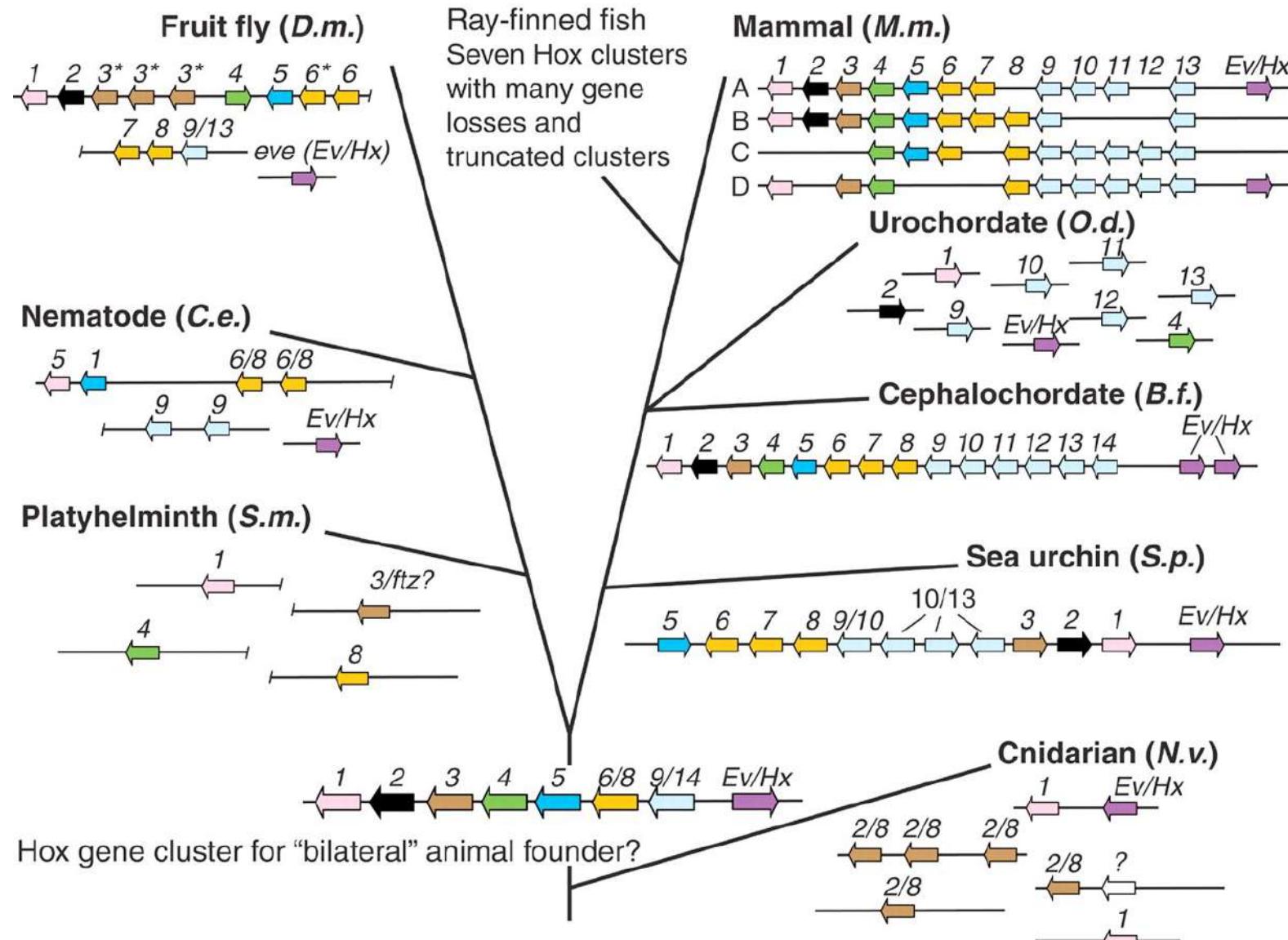
Genome duplication

- rare
- impacts whole genome

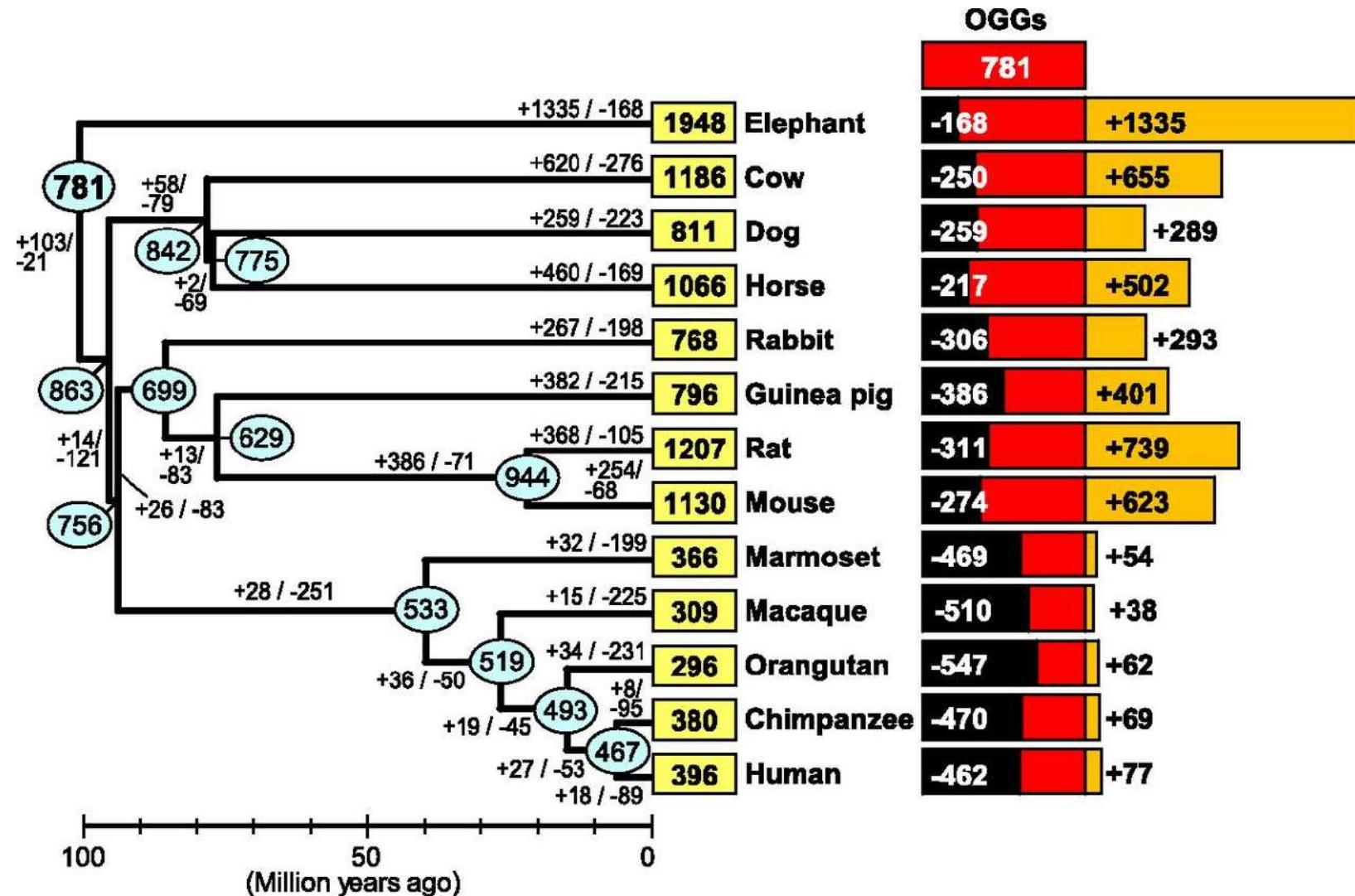
Discussion

What's a higher animal, how does it relate to gene or genome duplication?

Hox cluster history



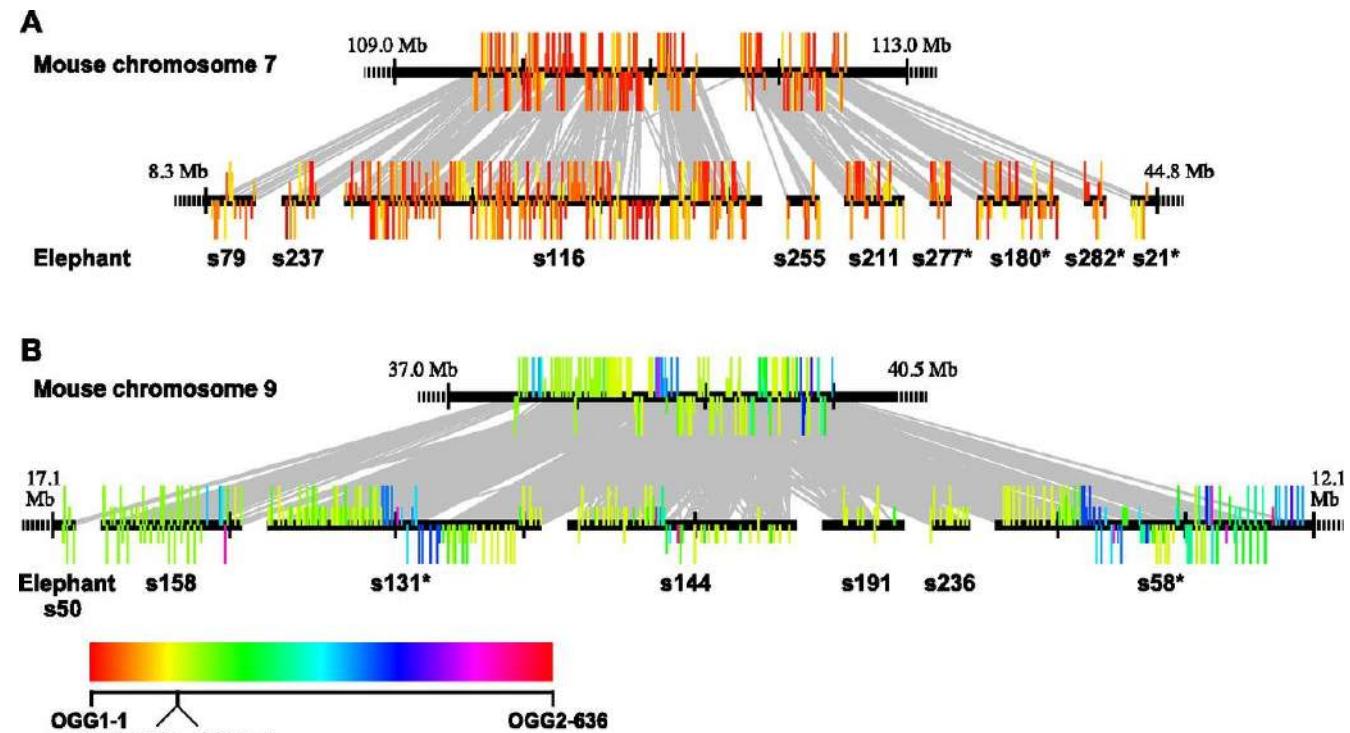
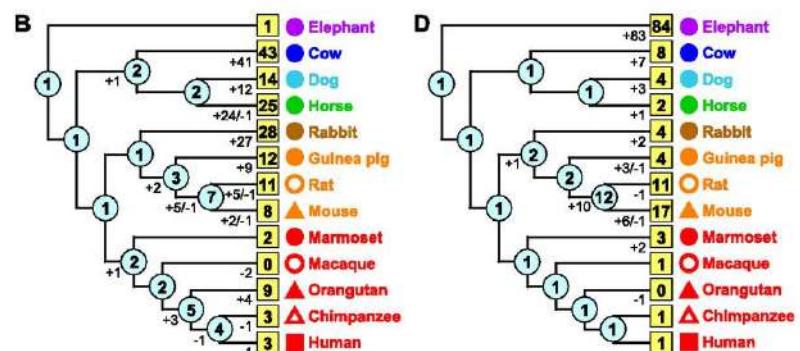
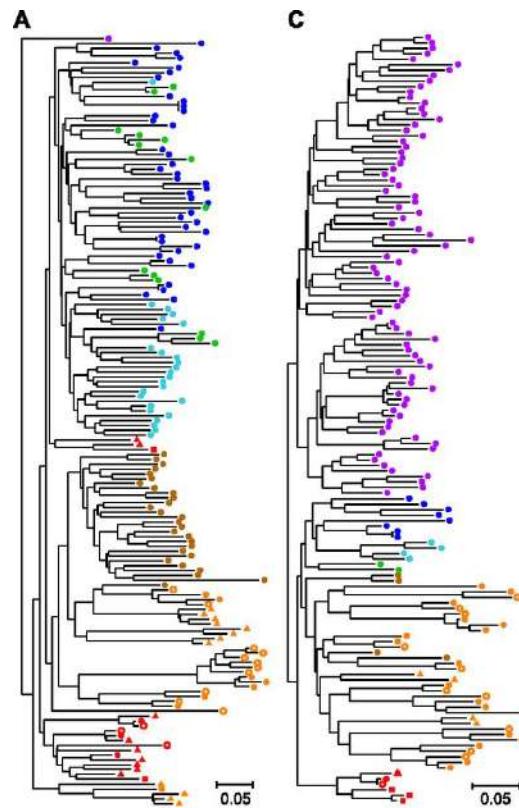
Mammal olfactory receptor genes



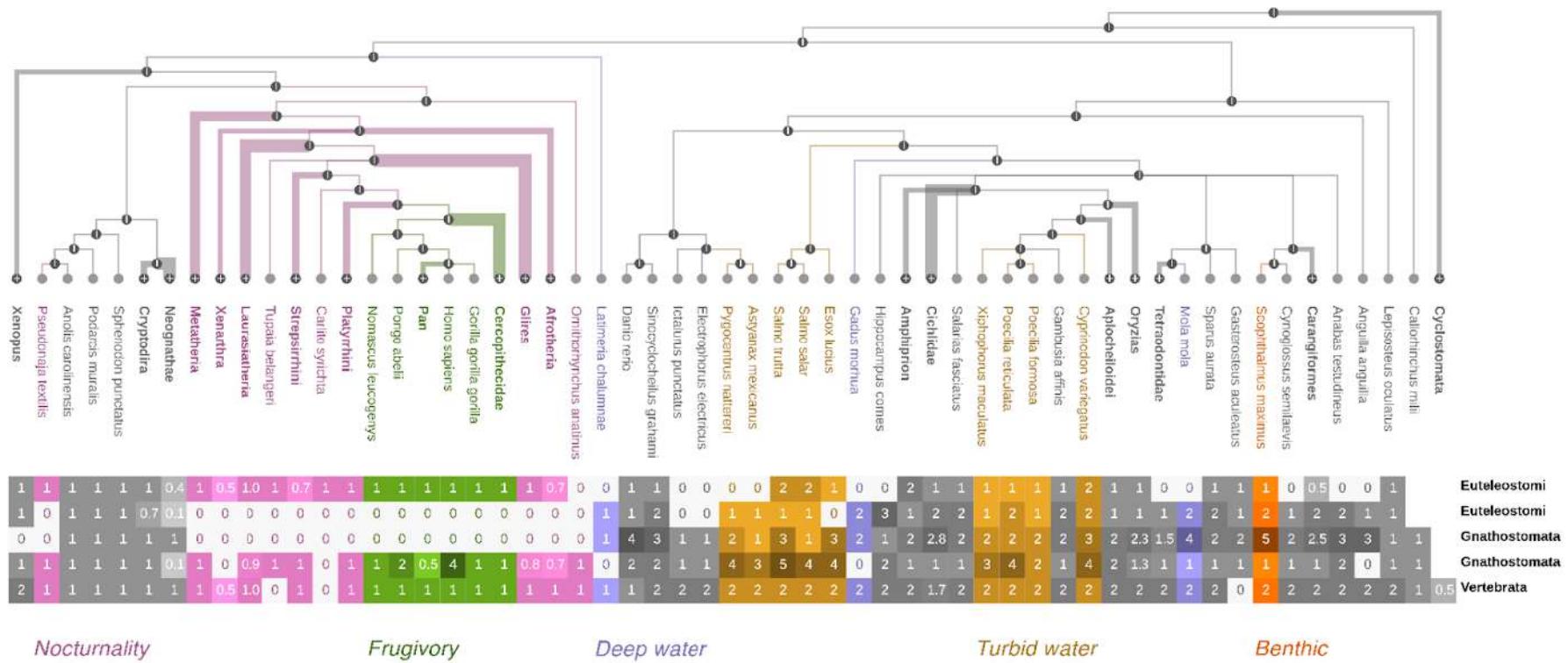
OGGs = ortholog gene groups

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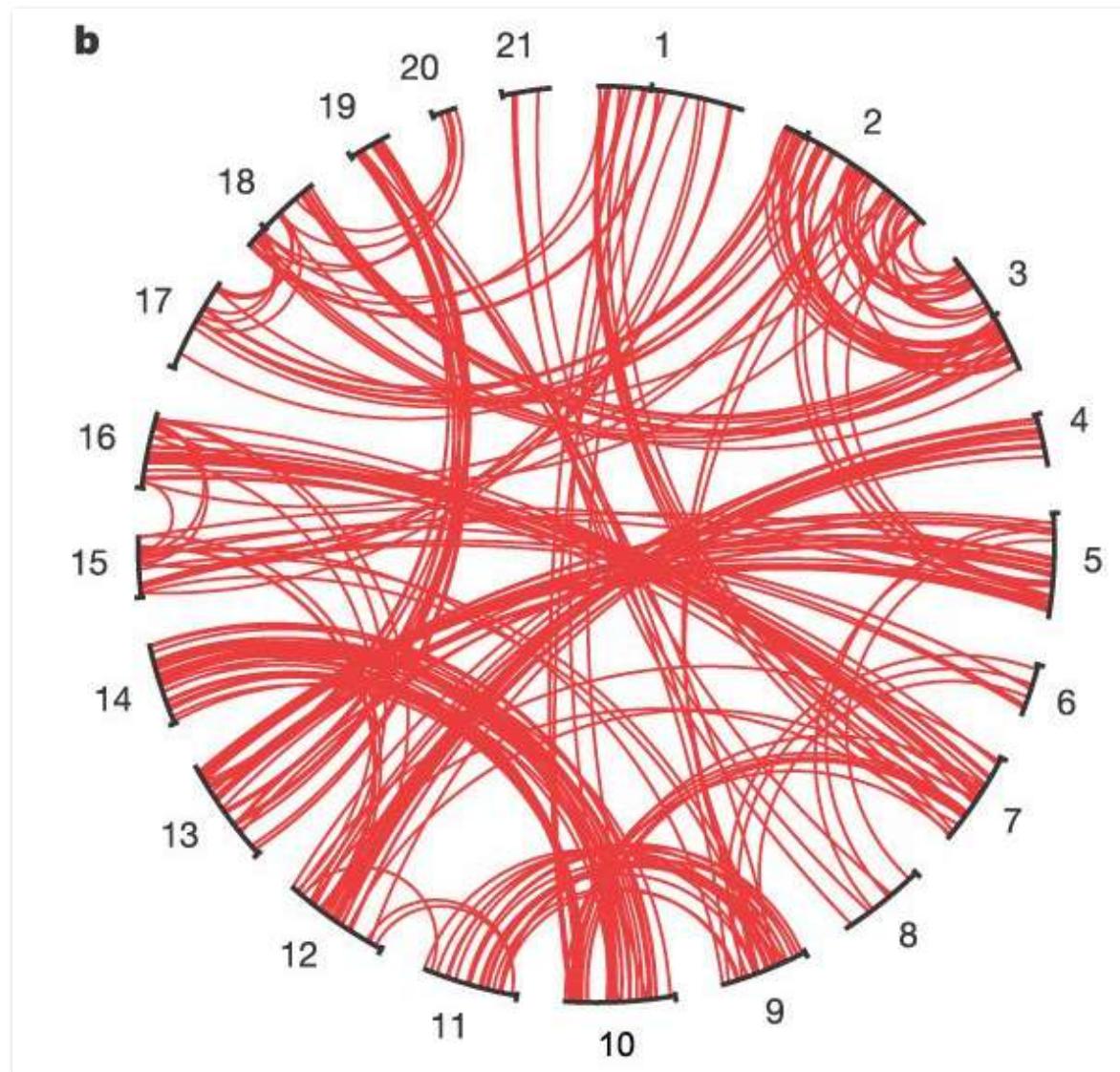
Elephant olfactory receptor genes



Visual opsin families in vertebrates

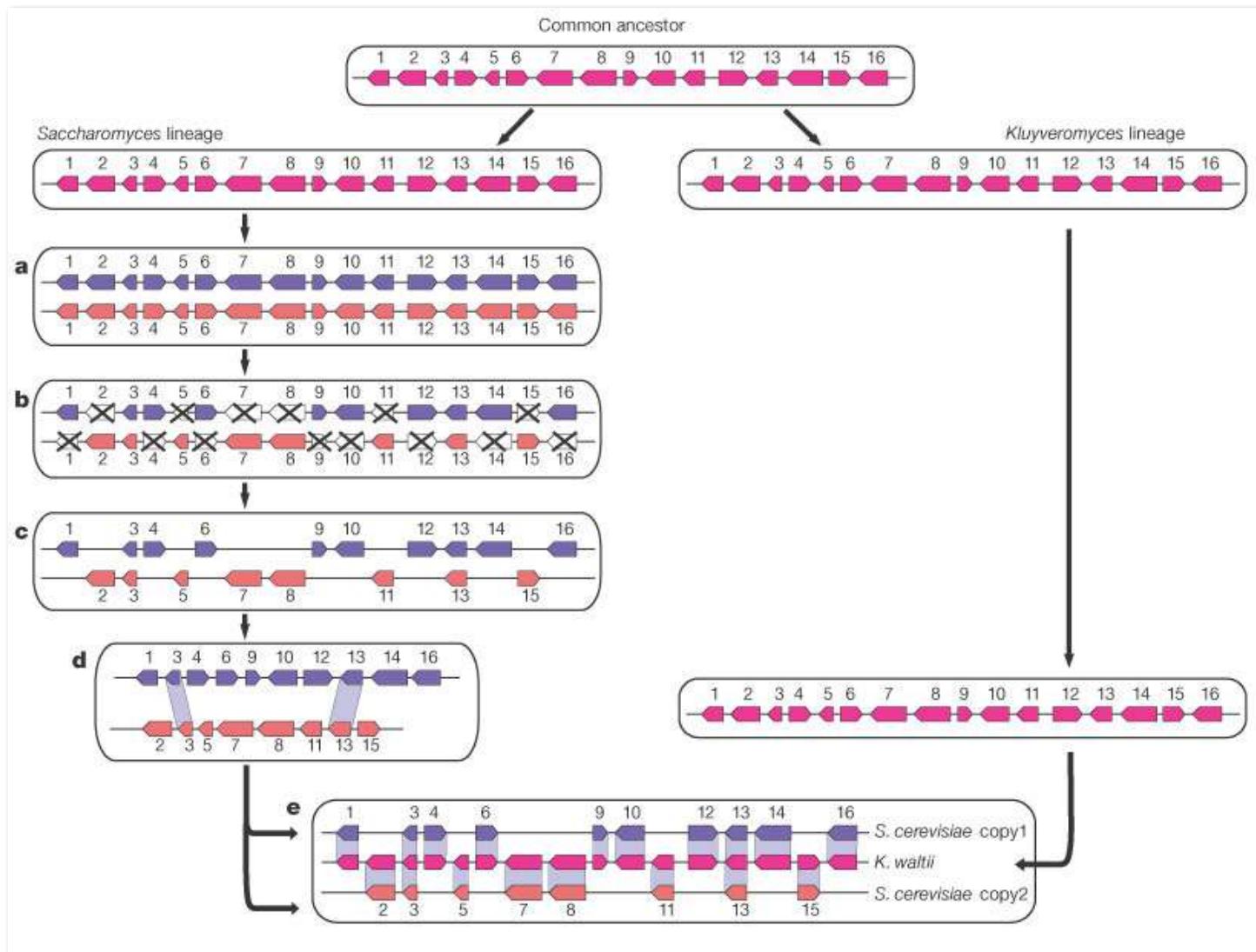


Genome duplication in teleost fishes

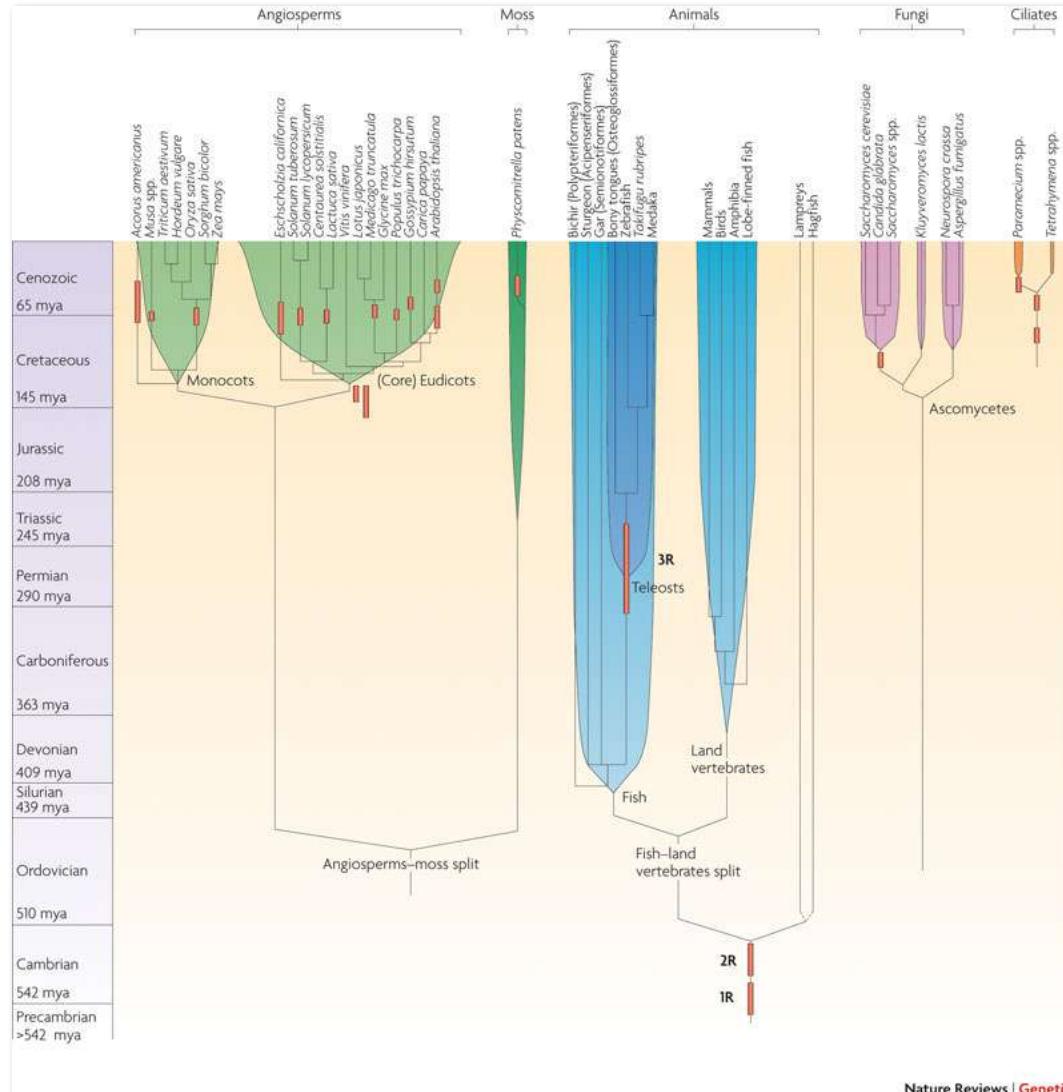


Global distribution of ancient duplicated genes ($K_s > 0.35$) in the *Tetraodon* genome

Whole genome duplication in yeast followed by gene loss

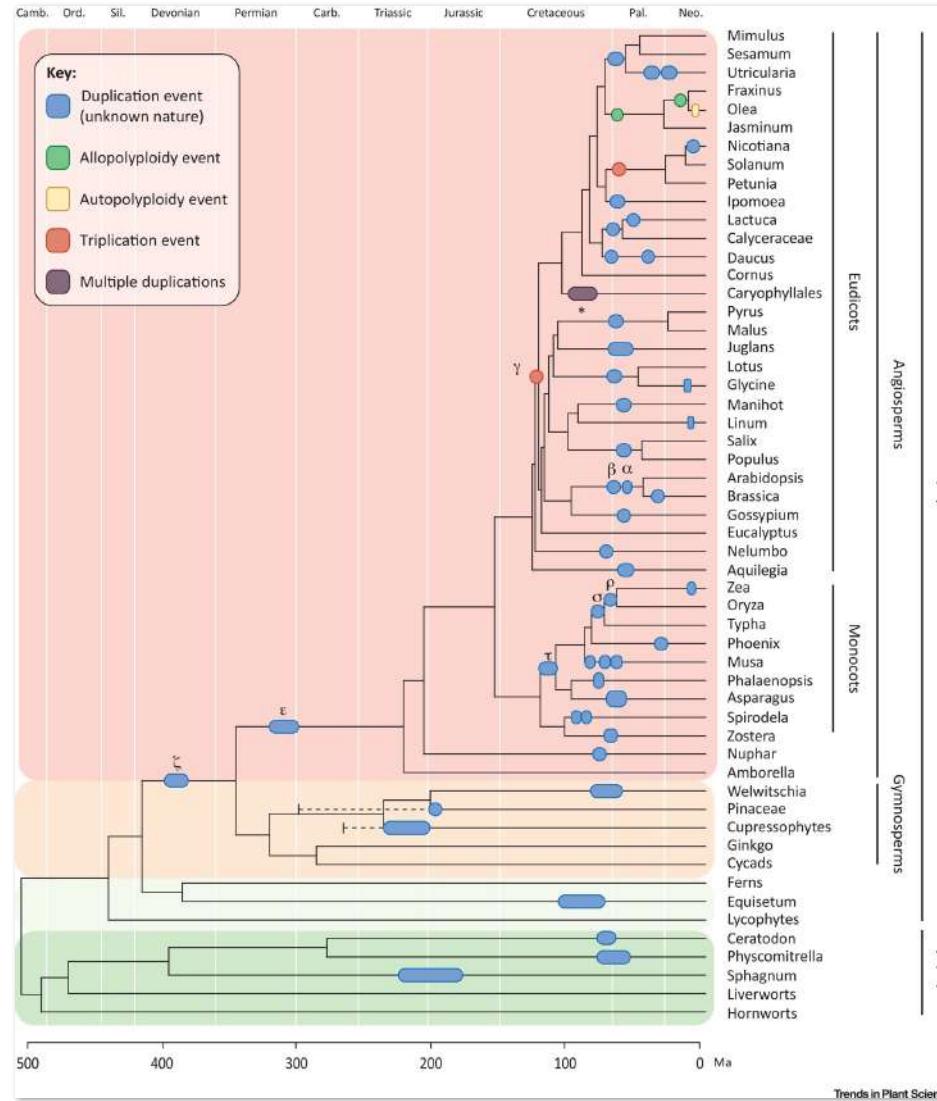


Genome duplications across eukaryotes

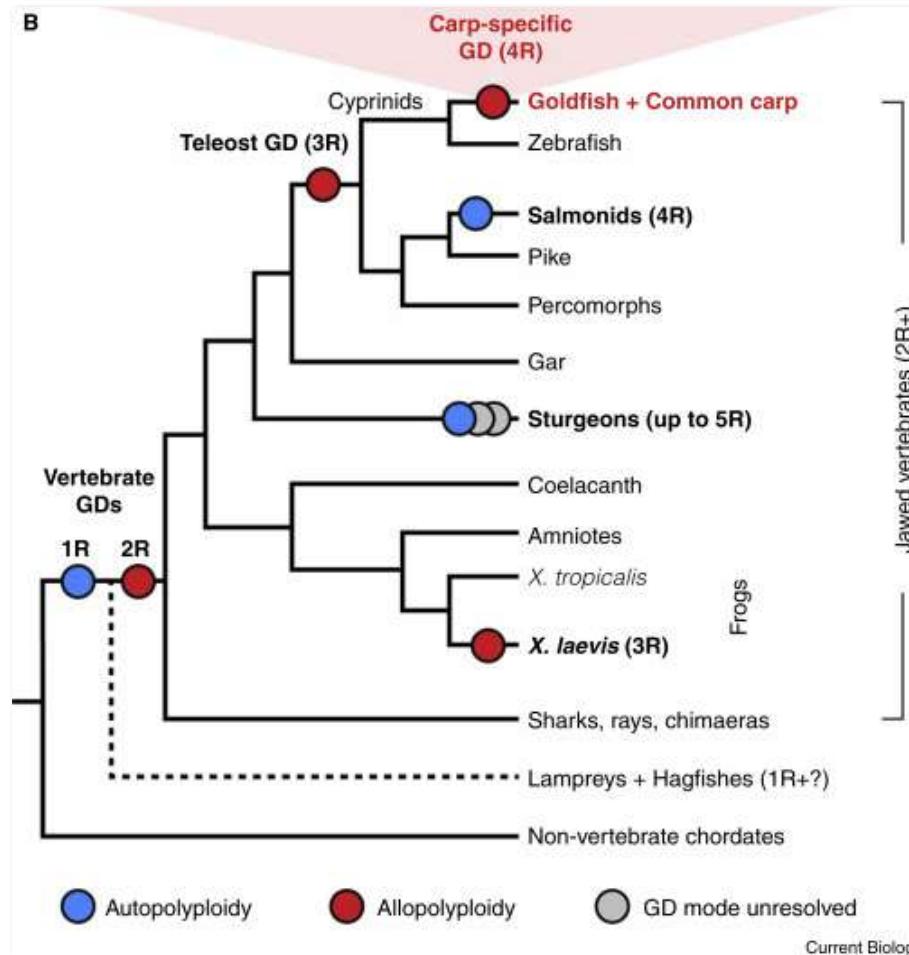


Nature Reviews | Genetics

Genome duplications in plants

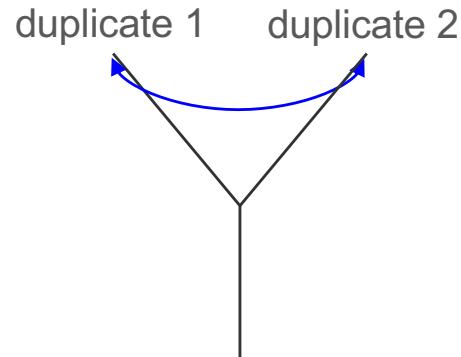


Genome duplications in vertebrates

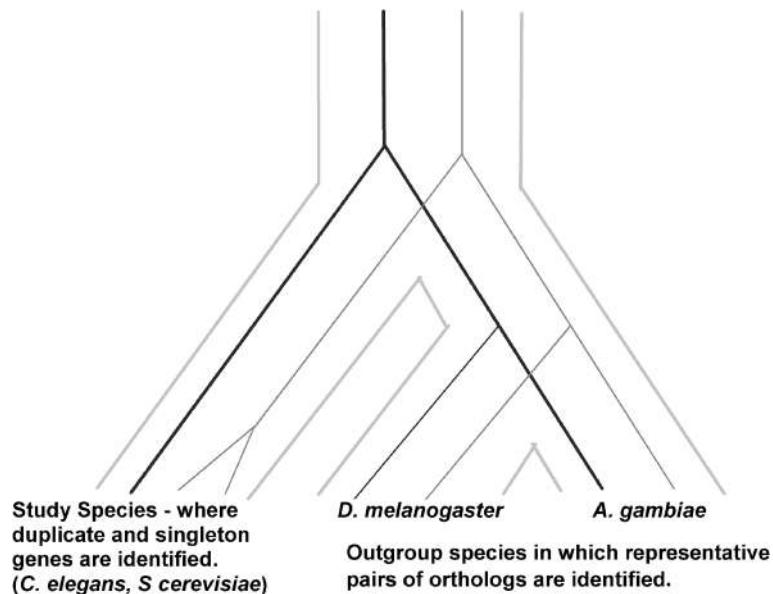


Evolution after duplication

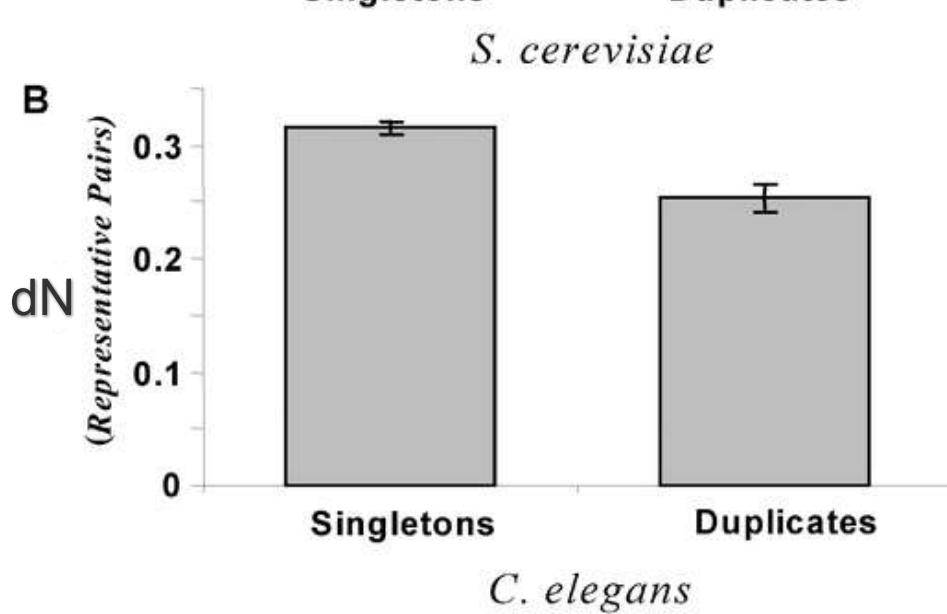
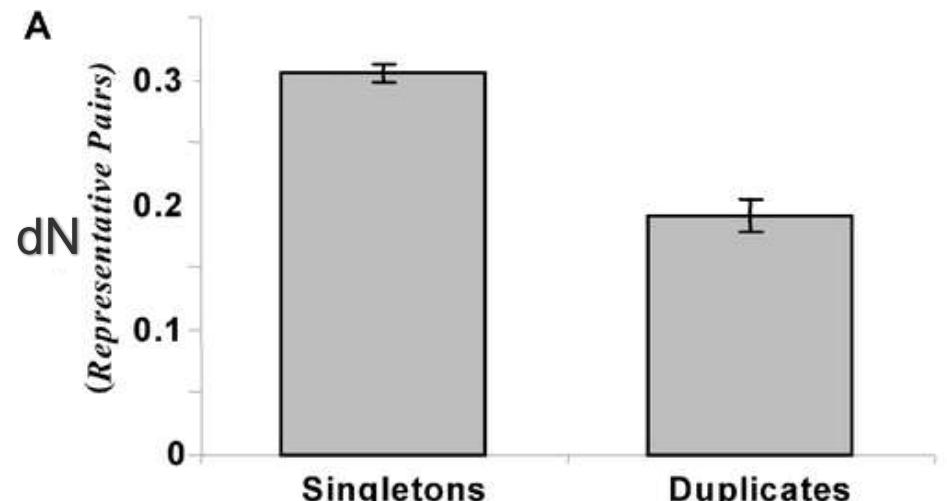
Biased retention of duplicate genes



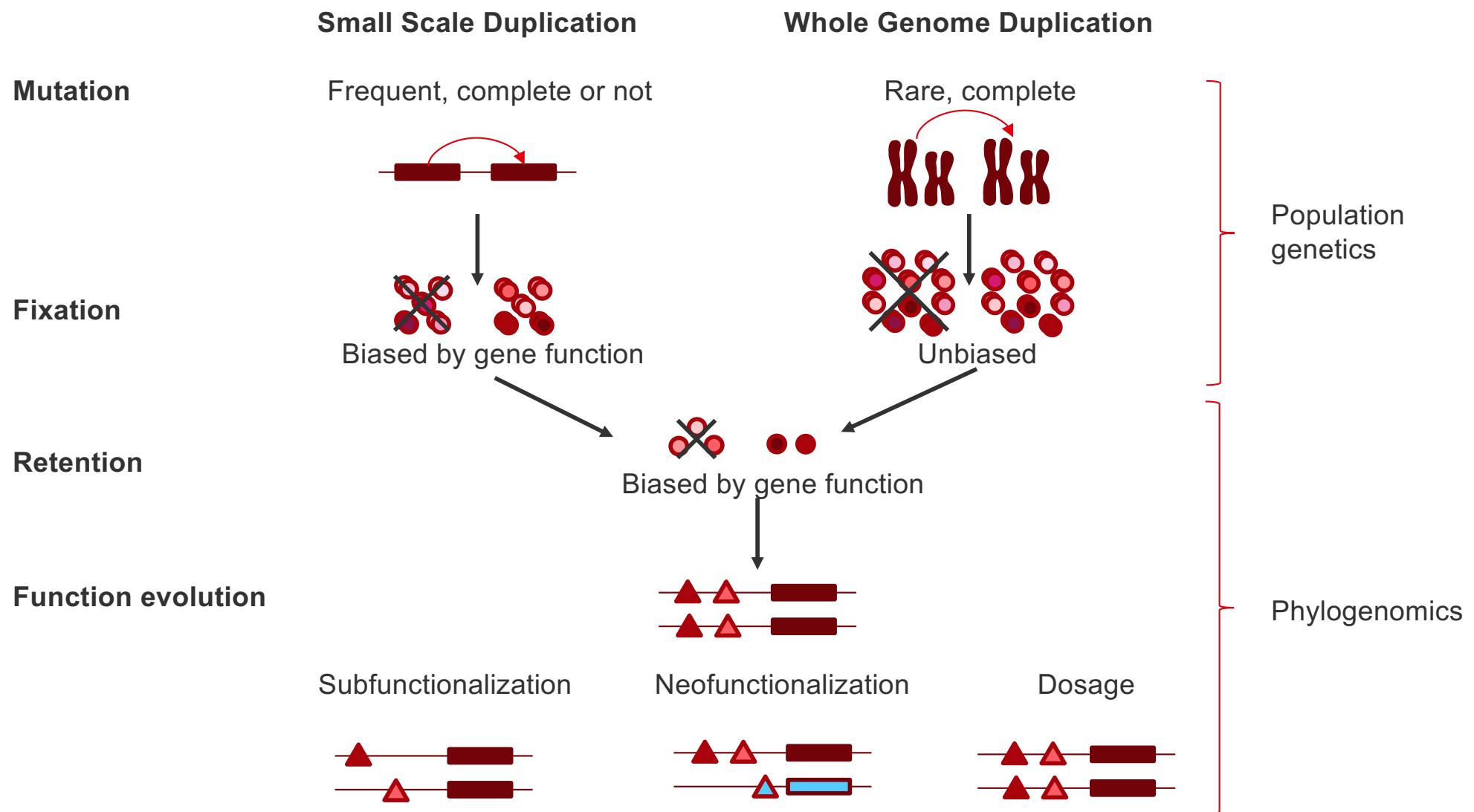
dN = original rate of gene
+ possible change due to duplication



dN Drosophila / Anopheles = independent of
changes after duplication in yeast or nematode

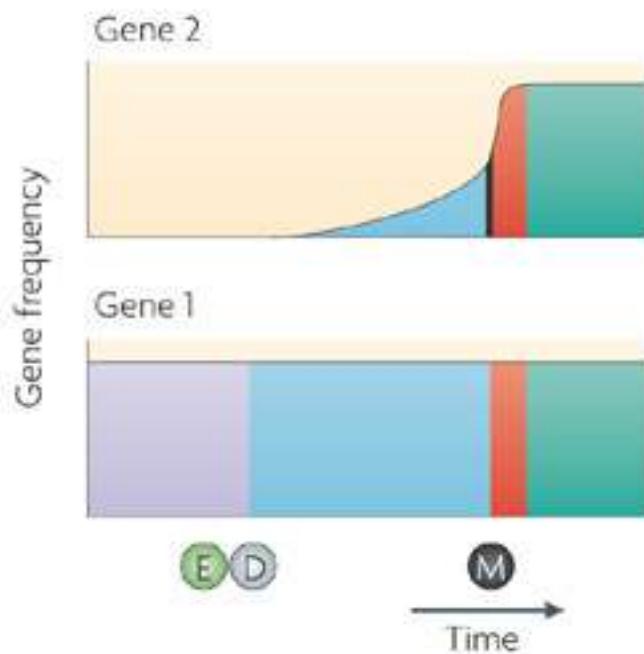


Impact of small- and large-scale duplications

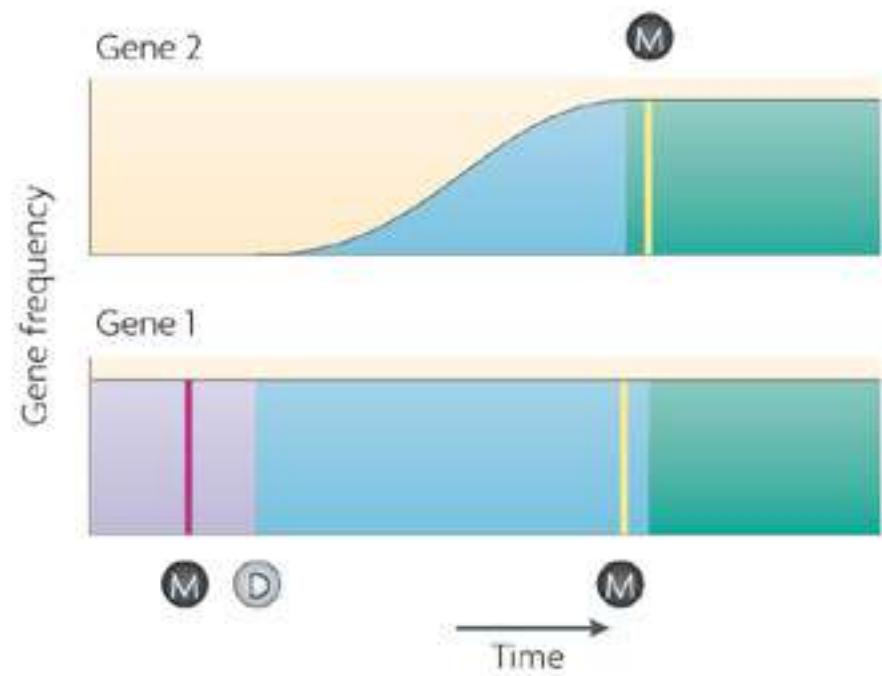


Selection and duplication

a N neofunctionalization

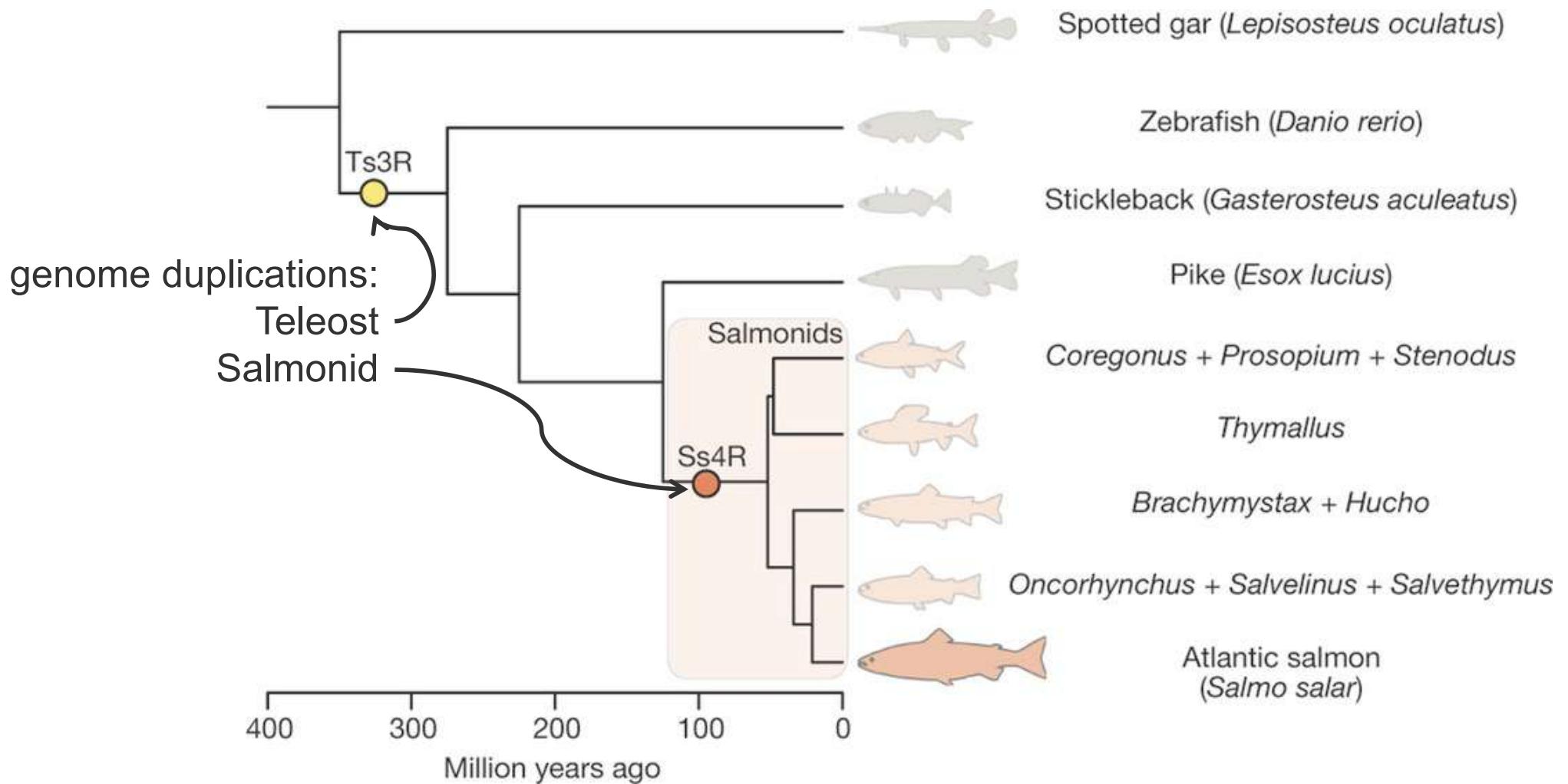


b c subfunctionalization

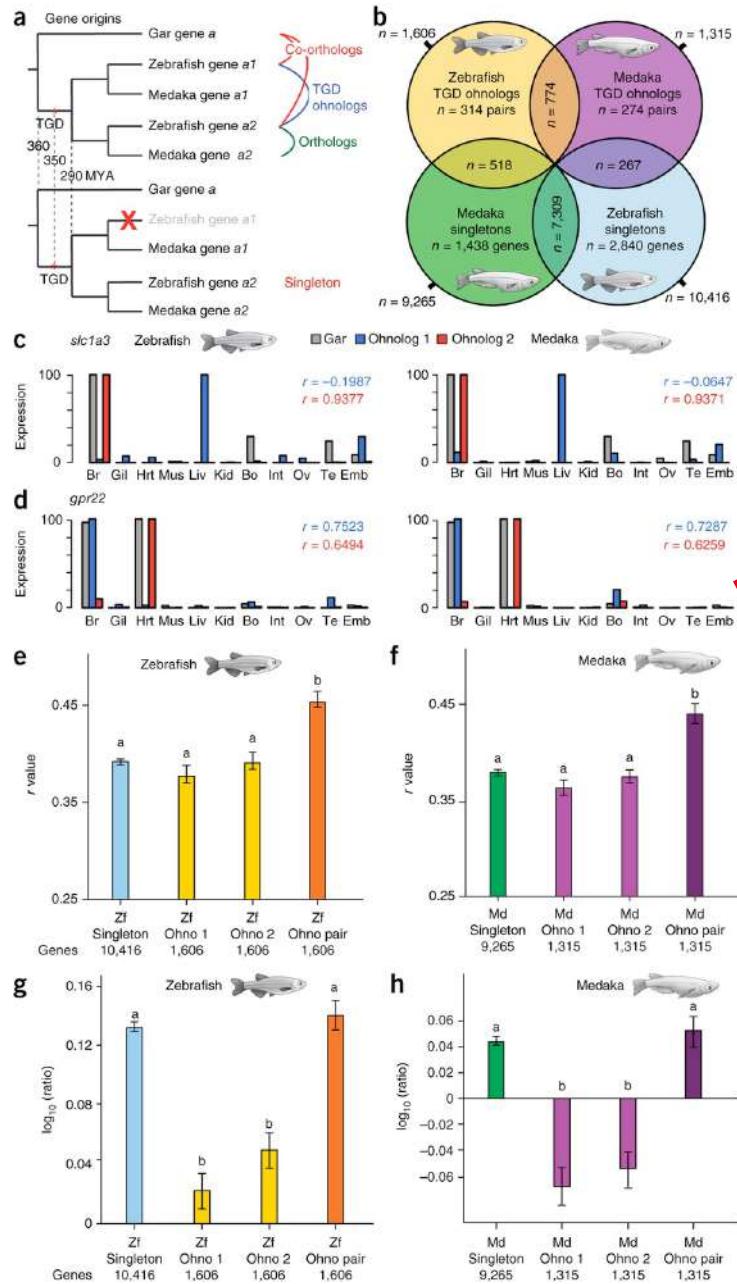


| | | | | |
|---------------|-----------------------|---------------------|------------------------------|----------------------------------|
| Genetic drift | Directional selection | Purifying selection | Appearance of novel function | Appearance of bi-functional gene |
| Mutation | Environmental change | Duplication | Loss-of-function mutation | Function-optimizing mutation |
| | | | | |

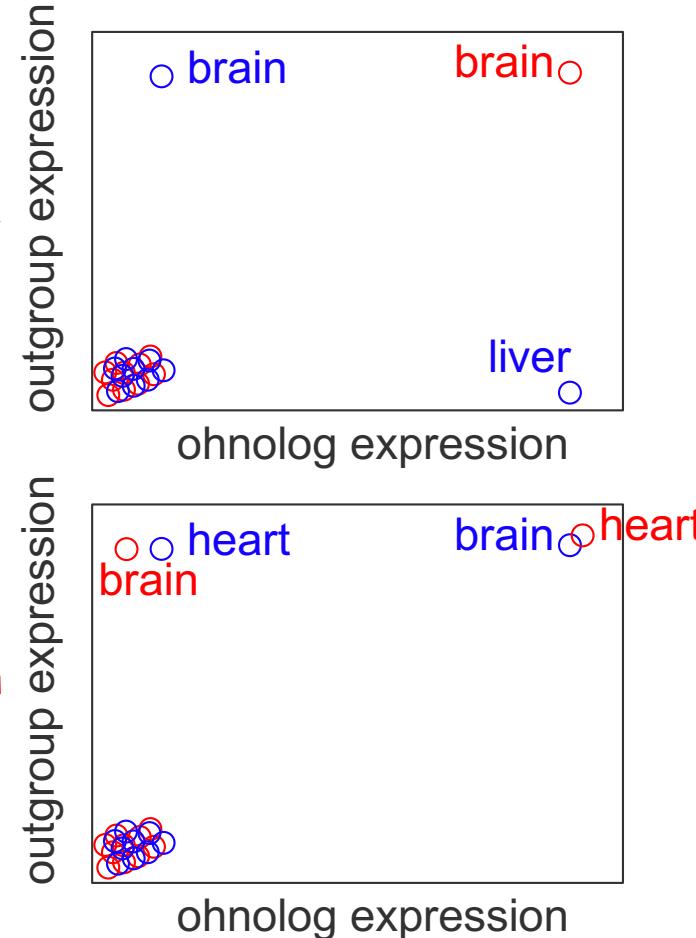
Evolution of expression after fish WGD



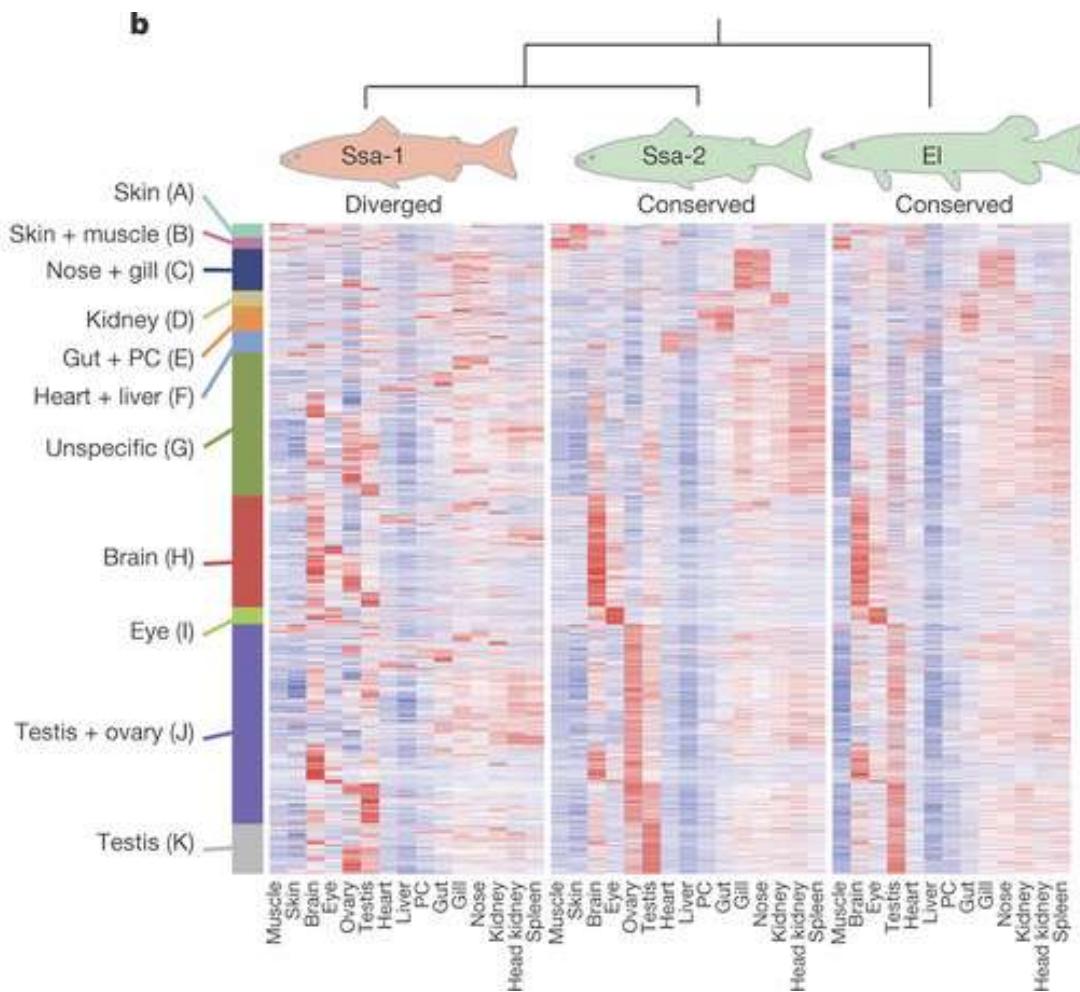
Support for sub-functionalization from teleosts



Comparison of expression:
RNA-seq in 10 organs or embryos of gar,
medaka and zebrafish



Support for neo-functionalization from salmonids



Comparison of expression:
RNA-seq in 13 organs or embryos of
salmon (*Ssa* = *Salmo salar*) and pike (*El*
= *Esox lucius*)

Evolution after genome duplication

- Source of information: expression
- Both sub- and neo-functionalization
- Many genes do not diverge significantly

Take home

Discussion

What do you take home from today?

**Diversity of genomes
makes sense in the
light of gene function**
