

Evolution of the Vertebrate Y RNA Cluster

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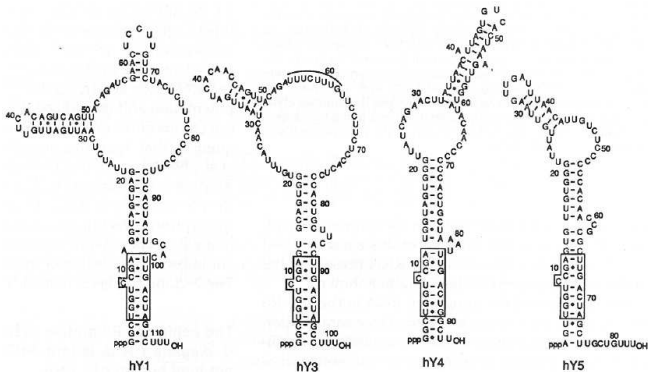
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Outline

- 1 Introduction
- 2 The Y RNA Cluster
- 3 Evolution
- 4 Discussion

What are Y RNAs?



Human Y RNAs: 4 paralogs located in a single cluster on Chr.

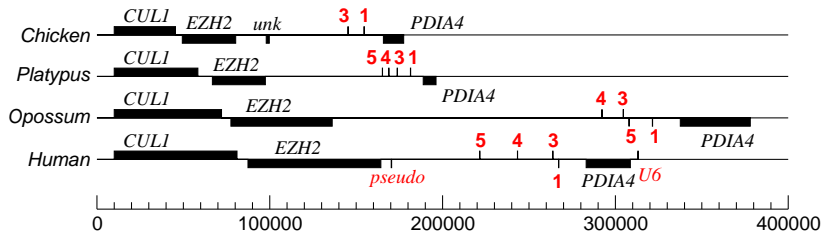
What are Y RNAs?

- Short non-protein-coding RNAs
- Transcribed by RNA polymerase III (like tRNAs, U6 RNA, vault RNAs, ...)
- Part of the Ro ribonucleoparticle
- Located in close proximity in humans, mouse, and xenopus
- direct role for DNA replication
- In human: many pseudogenes, apparently only a single functional cluster

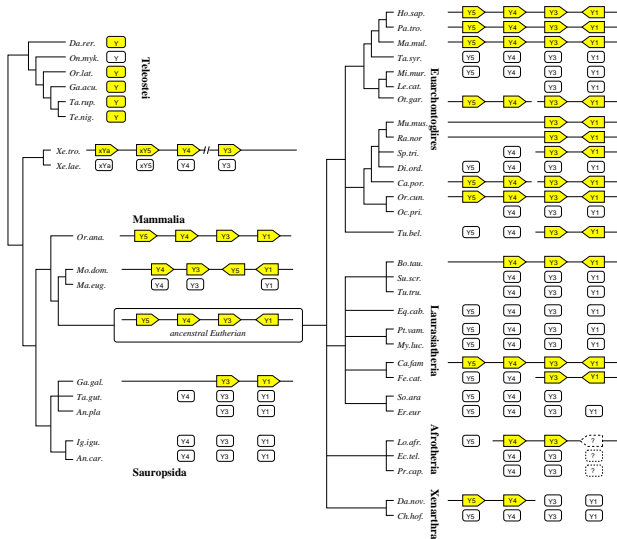
Previously known facts

- 4 Y RNAs in human but only two in mouse
- generally well conserved in mammals
- 2 Y RNAs in chicken
- 4 Y RNAs in xenopus, relation with mammalian ones unknown
- there are Y RNAs in teleost fishes

Genomic Location



Inventory



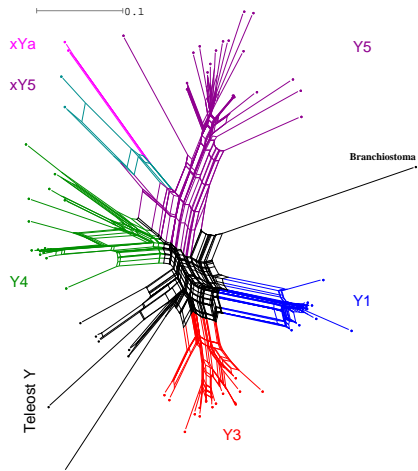
Deviant Y1 in Afrotheria

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Hs_Y1  GG-GGGGGAGGAGTGAATAAUAAAGATGCAAGCAGTACAGATGAAACCTGGTGAATGGCCCCTCACTAATGAACTGACTAGCT-- 110
Lo_afr-Y1-p  GGAGGGGGAGGAGTGAATTTCTGGGAG--TTCTCAGCCAACTACAGGTGAACTTGGTGA-----CCCGCTCACTGATGATGACTGGCTTT 102
Md_Y1  GG-GGGGGAGGAGTGAATAAAAAGATGCAAGCAGTACAGATGAAACCTGGTGAATGGCCCCTCACTAATGAACTGACTAGCTTTT 113
ruler  1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....

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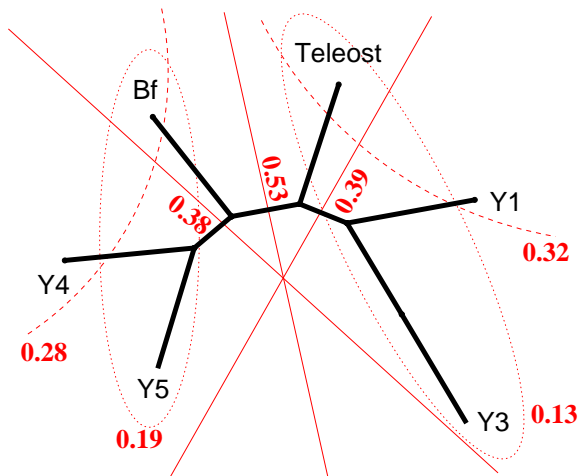

Neighbor Net



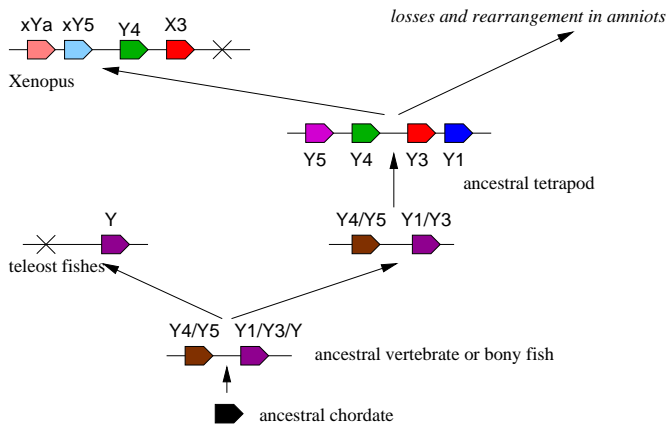
Splits between groups

- Assume that Y1, Y3, Y4, Y5 (incl xY5 and xYa), and teleost Y are monophyletic
- Choose one sequence from each of these 5 groups plus the Amphioxus sequence
- For each combination compute a neighborjoining tree
- Extract the three non trivial splits from this tree
- Record the most frequently occurring splits
This defines the *strict consensus tree* from the three most frequent splits

Splits between groups



Summary: The most likely szenario



Open Questions

- Y RNAs have also been reported in *C. elegans* and in the bacterium *Deinococcus radiodurans*
- So far, no successful search for Y RNAs in flies, basal deuterostomes, plants, etc, although the Ro protein is evolutionarily old

Acknowledgements

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