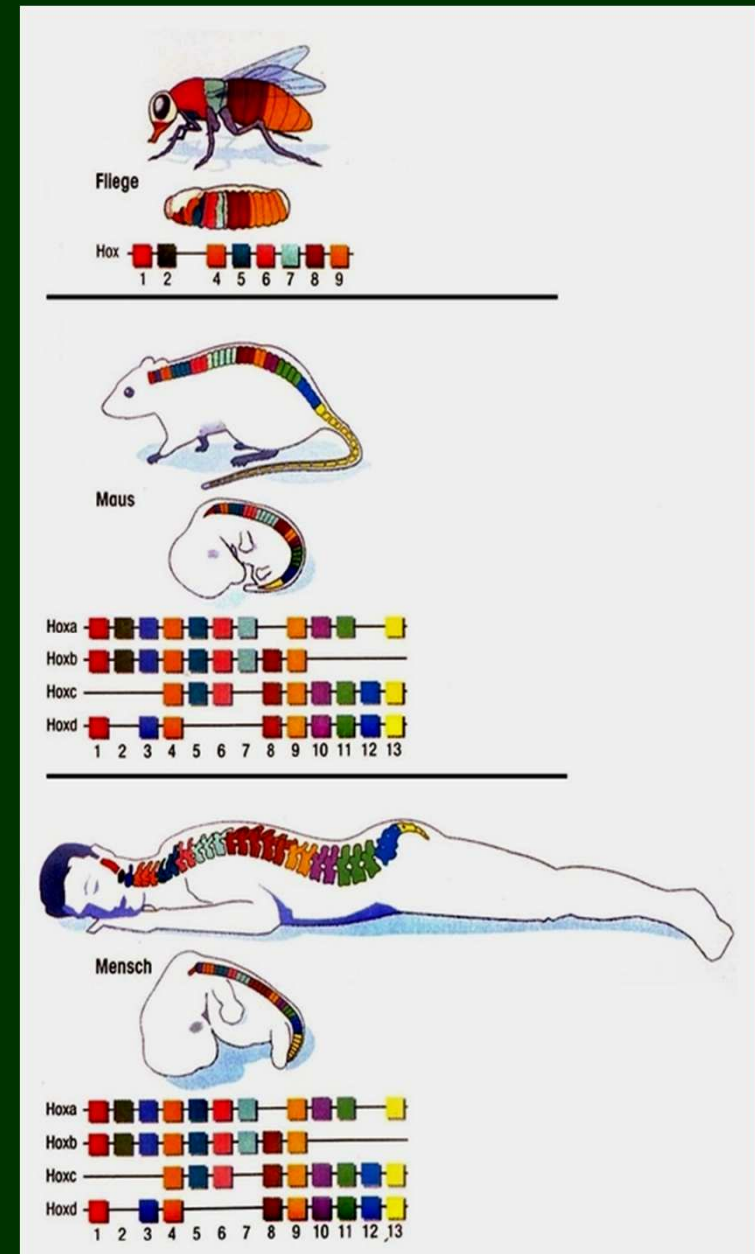


Bilateria: Xenacoelomorpha a Chaetognatha

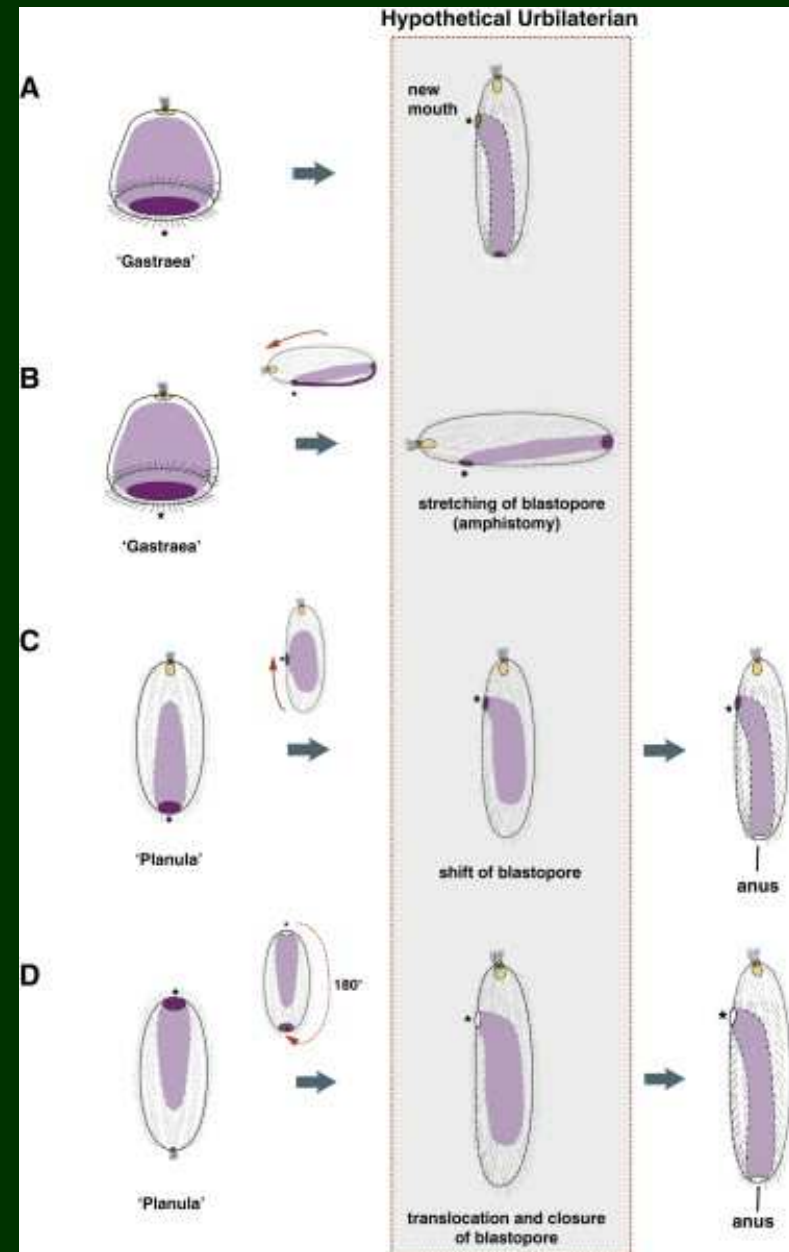


Bilateria

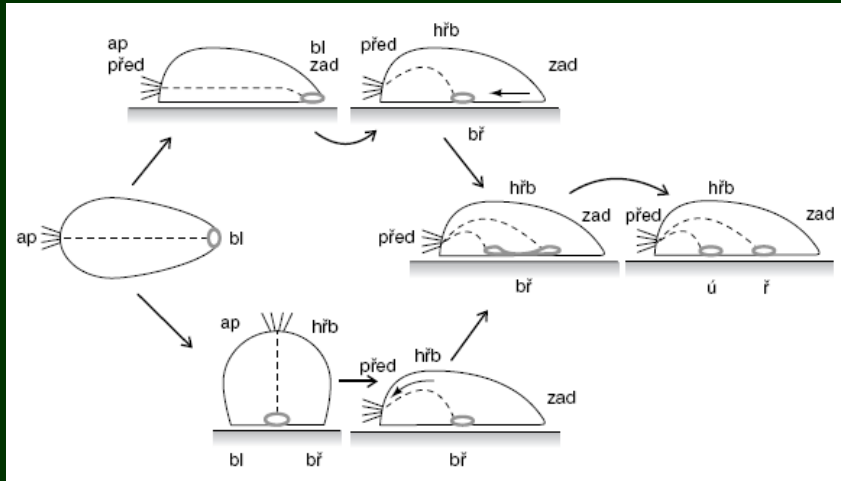
- dvě tělní osy: antero-posteriorní a dorso-ventrální
- *Hox* komplex
- trubcovitá trávicí soustava
- obvykle mnohobíčíkaté buňky epitelů
- céloom, cévní soustava, protonefridie



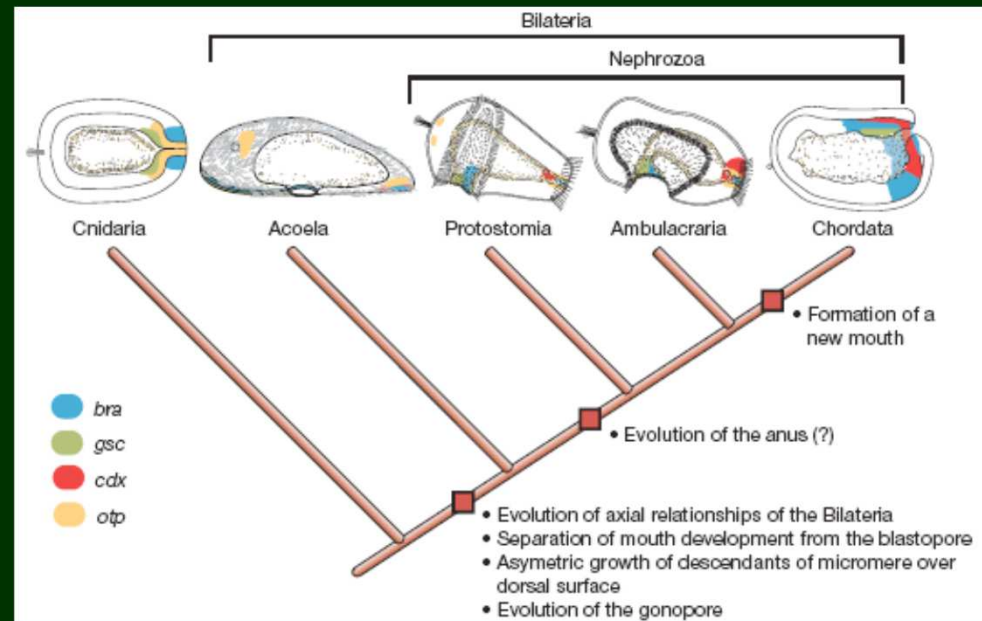
Vznik bilaterality



Vznik bilaterality

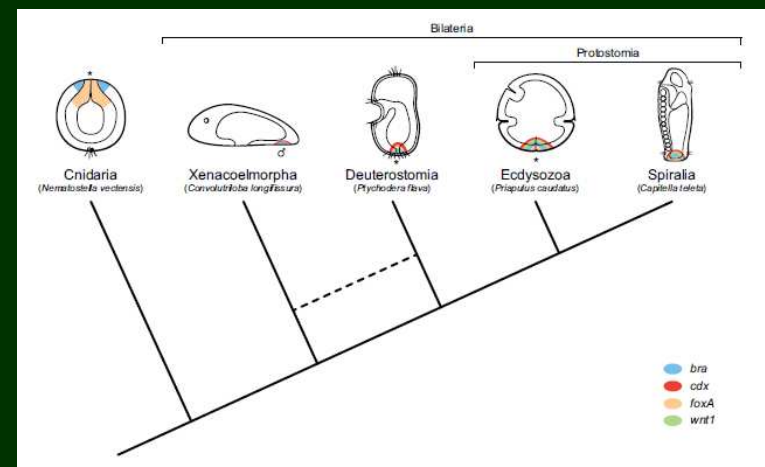
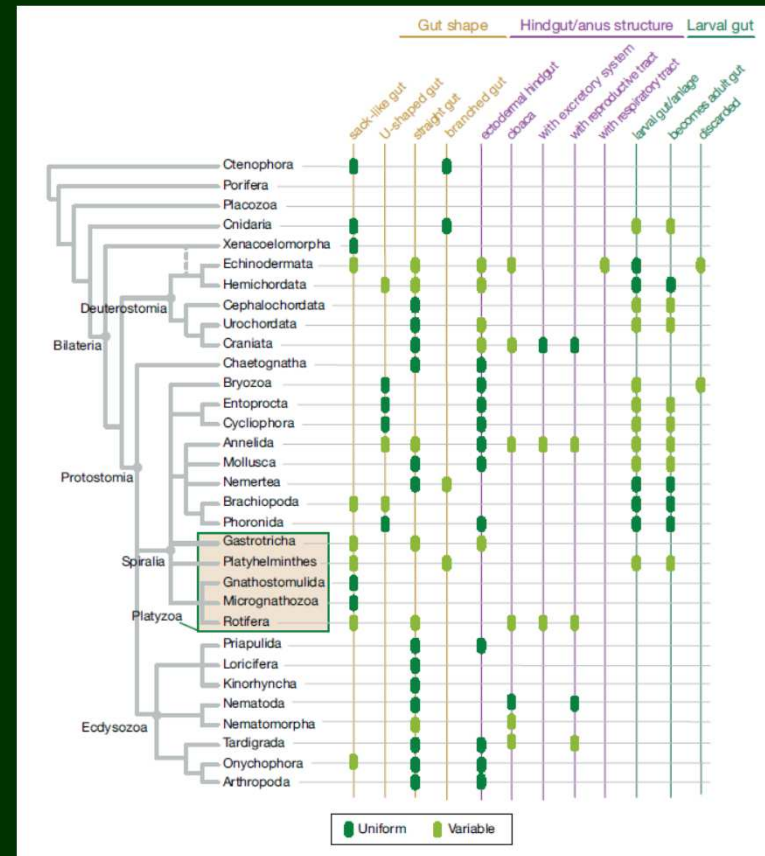


- **Acoela**: ústní otvor nevzniká z blastopóru (jako u mnoha protostomií a všech deuterostomií)
- gen *bra* není spojený s vývojem ústního otvoru (řitní otvor chybí), ale s gonopórem
- ústní otvor žahavců a acelomorf ~ ústní otvor bilaterií, řitní otvor vzniká nezávisle (endoderm + gonodukt)

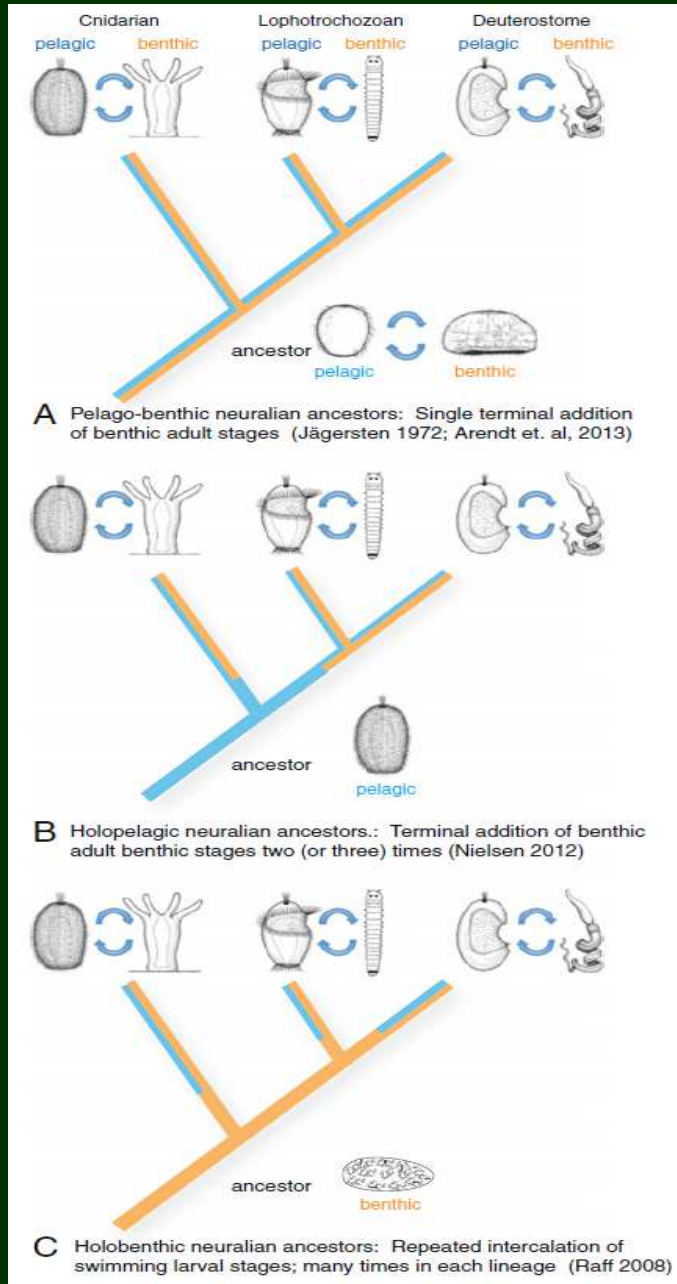


Trávicí trubice (řitní otvor)

- ústní otvor
acelomorf odpovídá
ústnímu otvoru
žahavců a nefrozoí
(*brachyury* +
gooseoid)
- řitní otvor ~ samčí
gonopór (Acoela)



Evolve životního cyklu



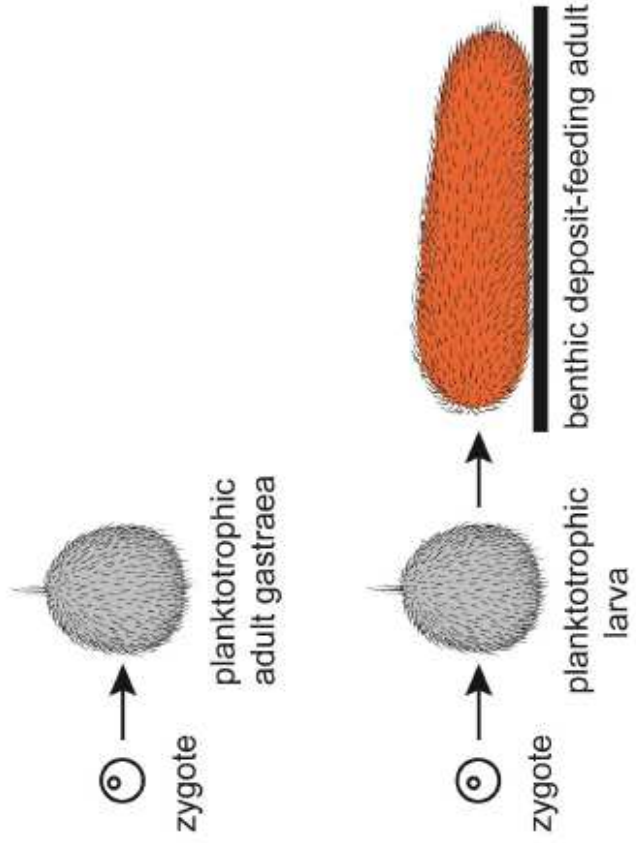
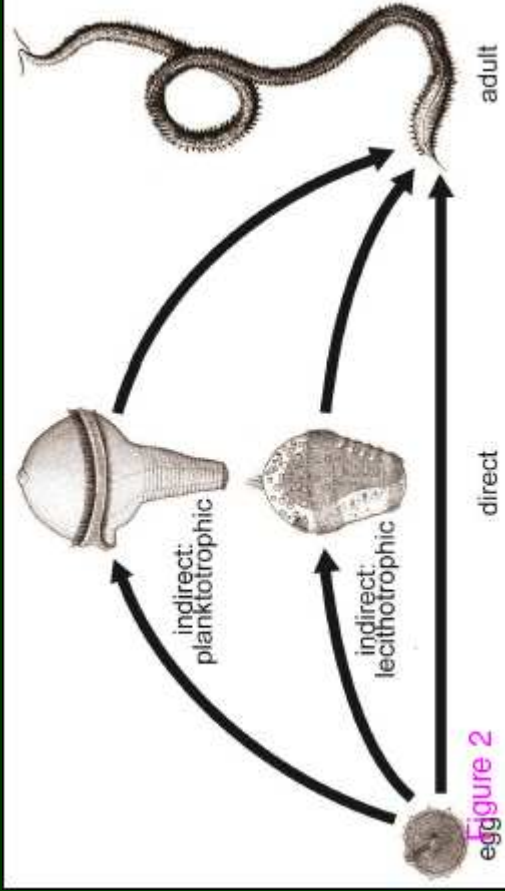
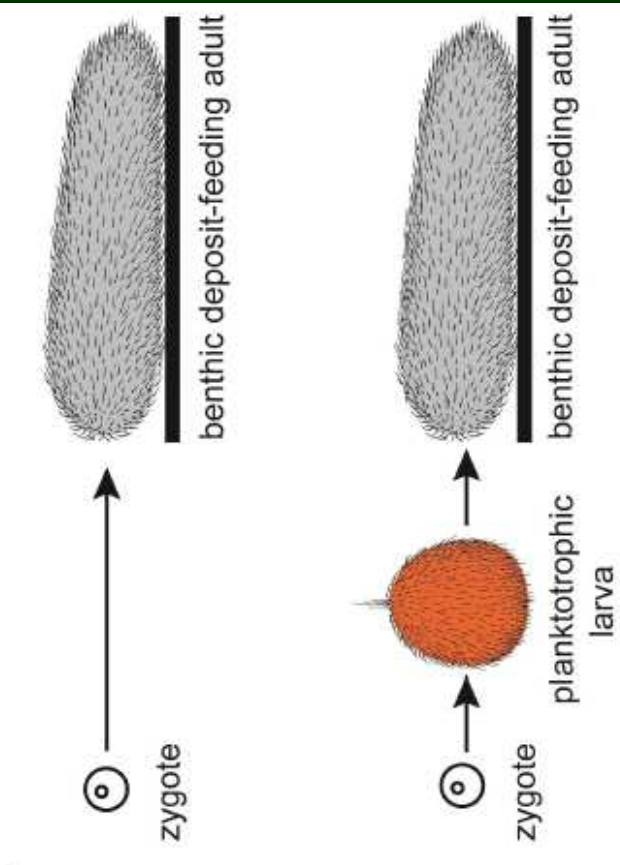
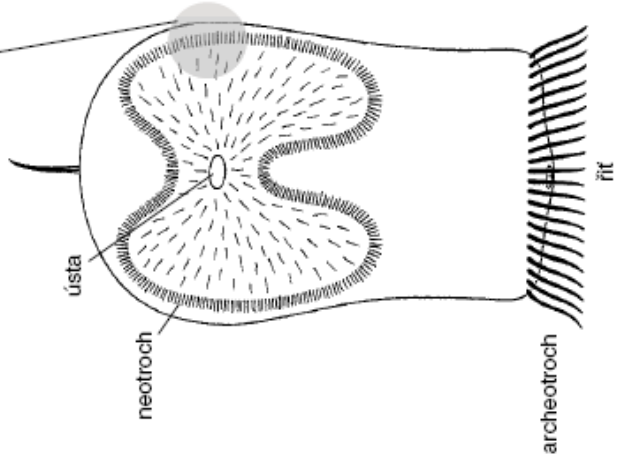
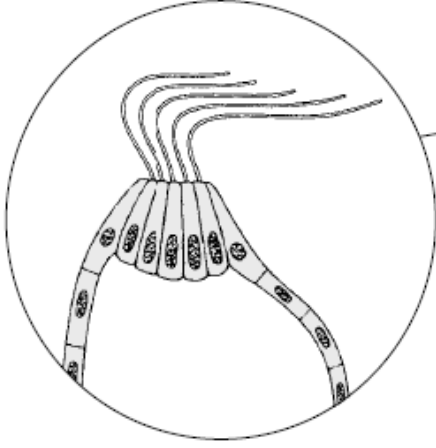


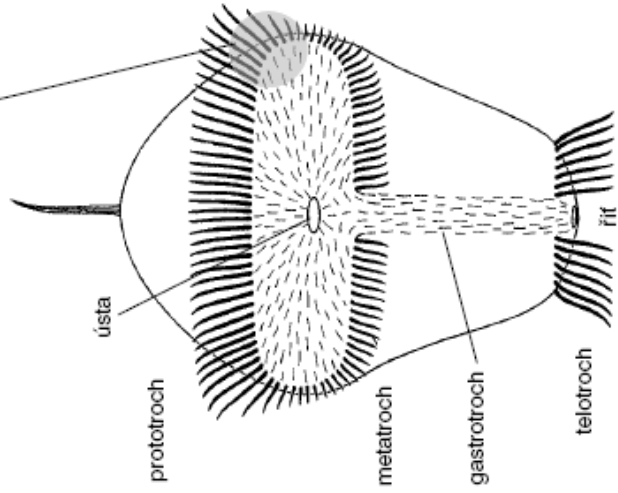
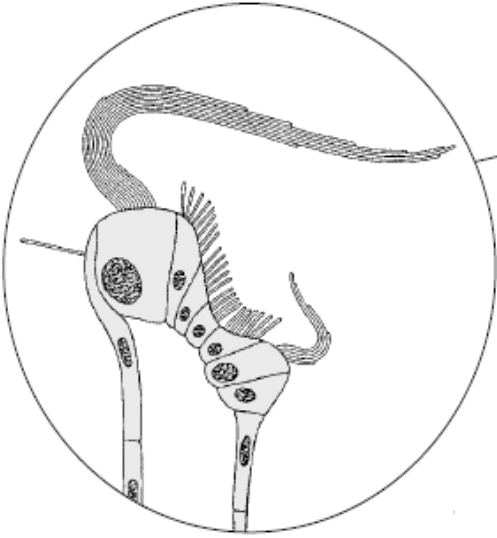
Figure 1 Terminal addition theory



Intercalation theory

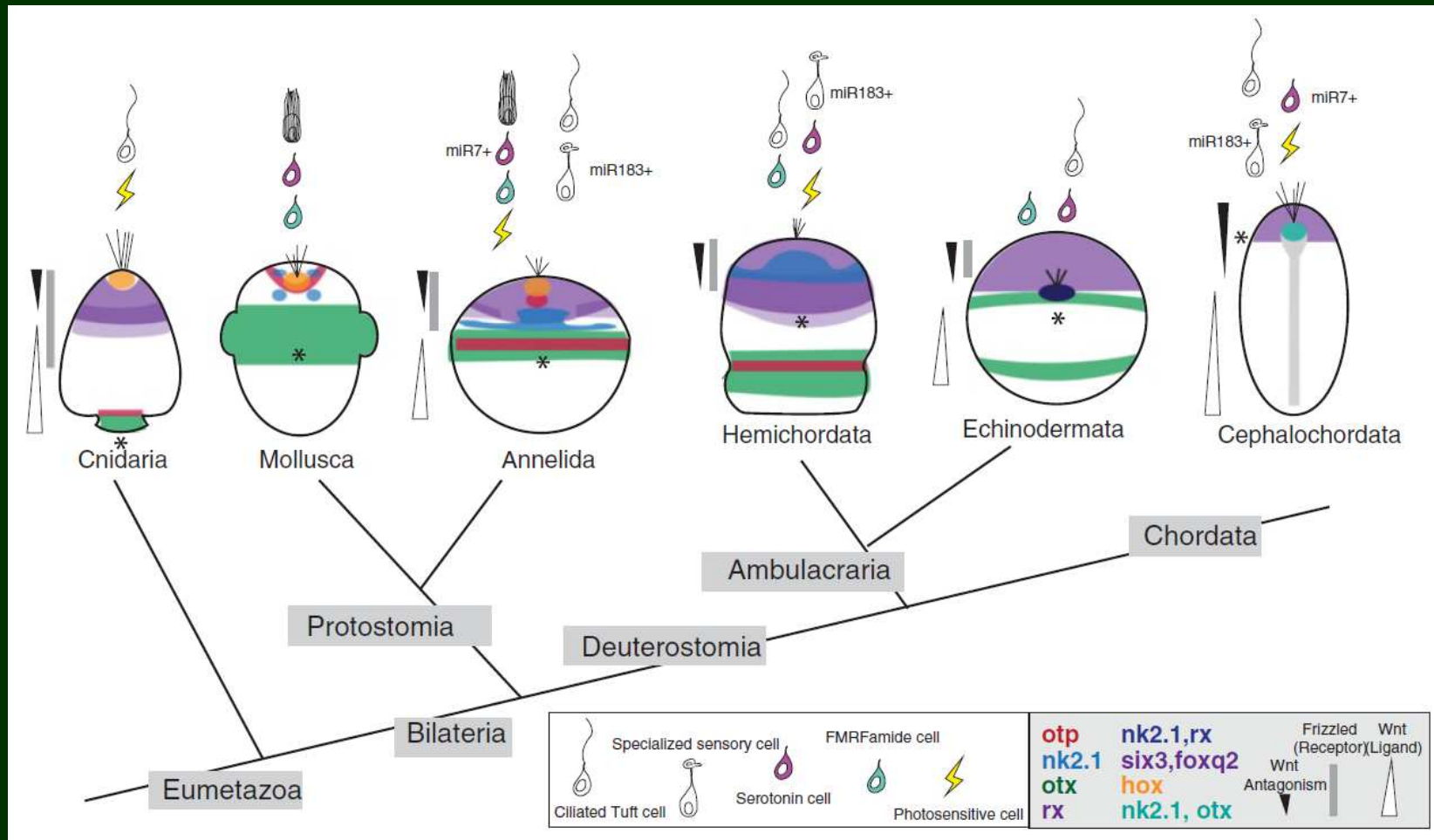


dipleurulov larva

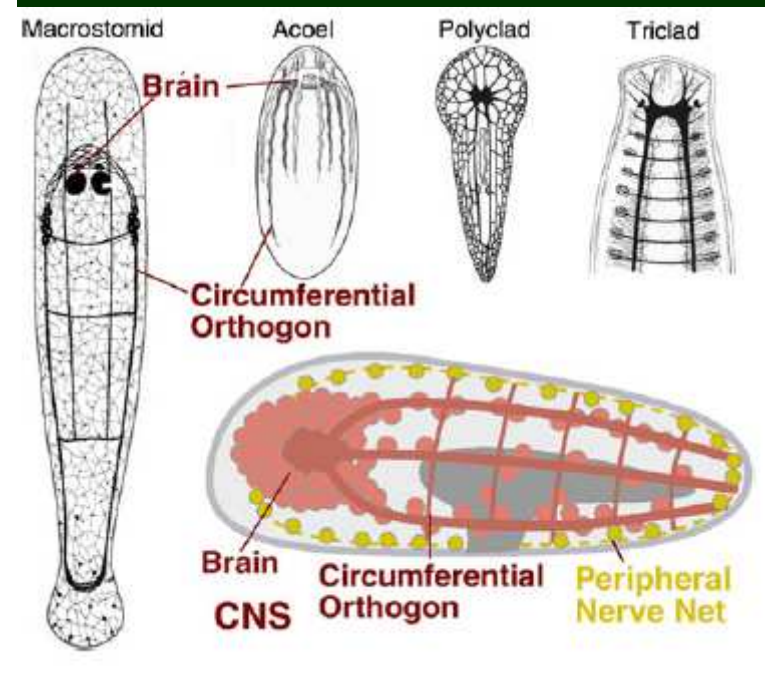
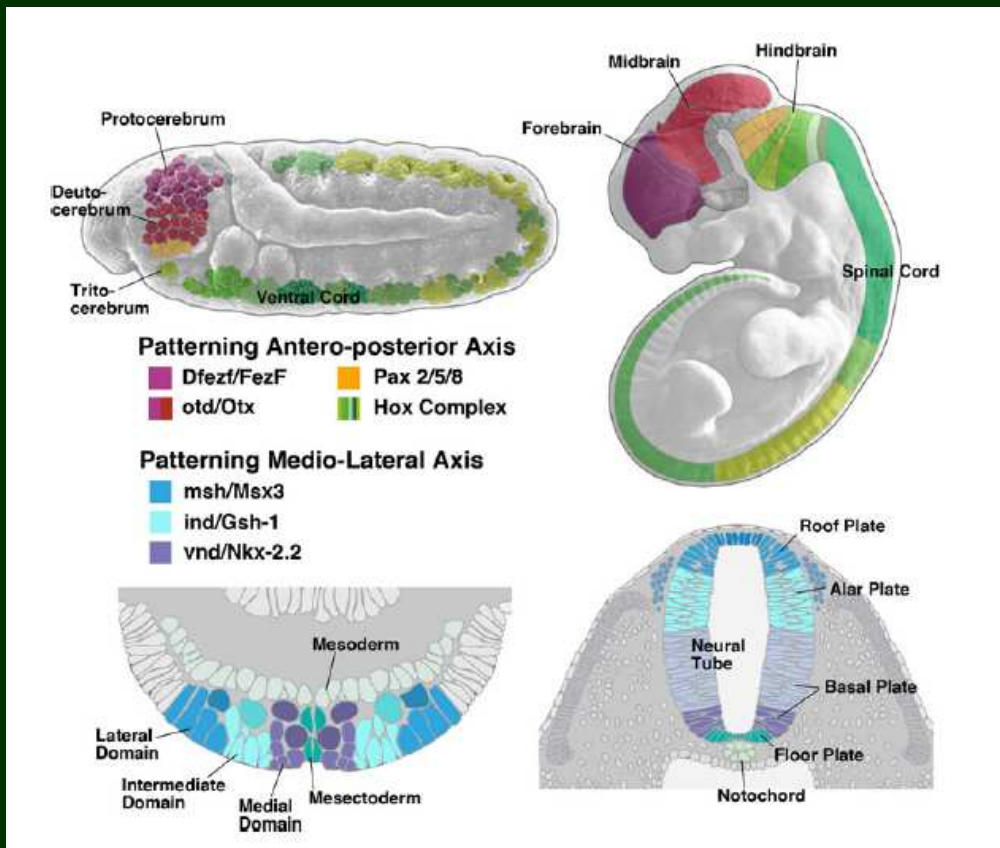


trochoforov larva

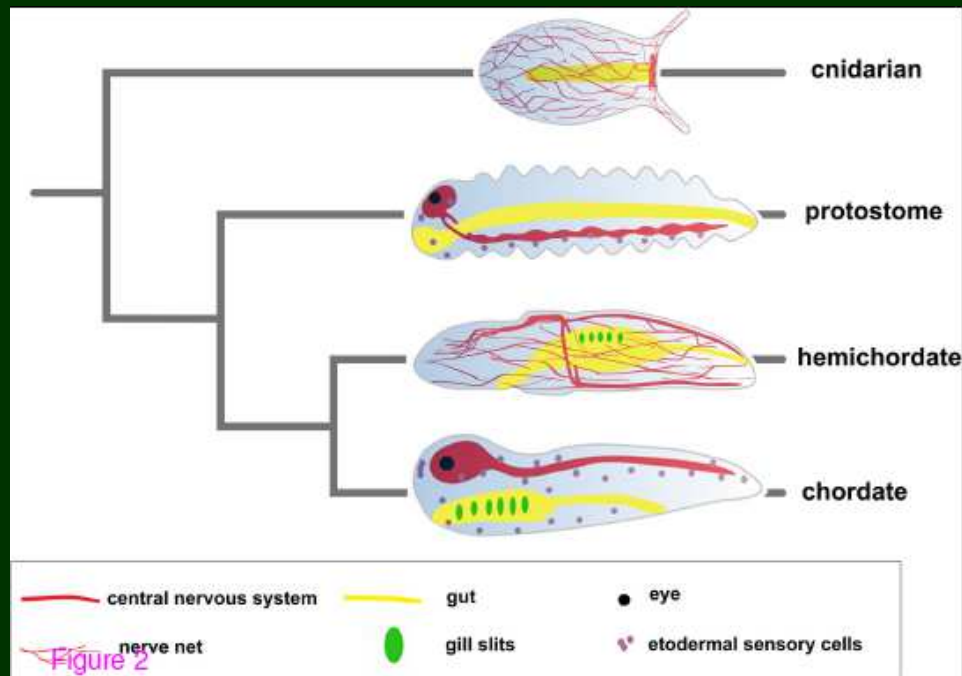
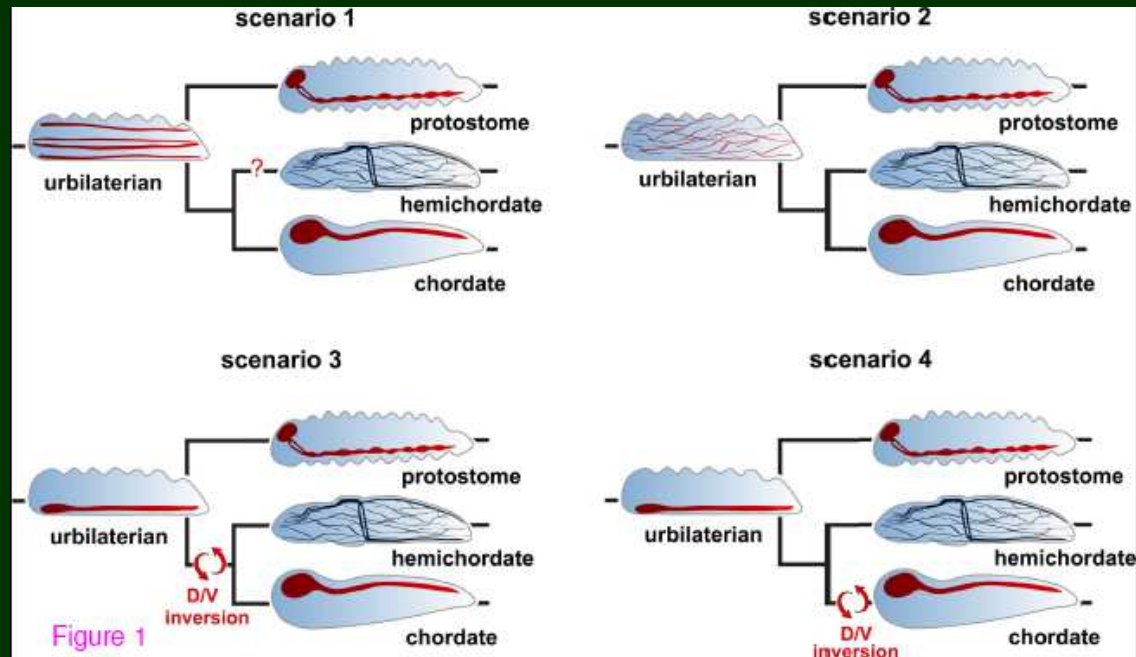
Apikální orgán a NS: homologie primárních larev?



Nervová soustava bilaterií

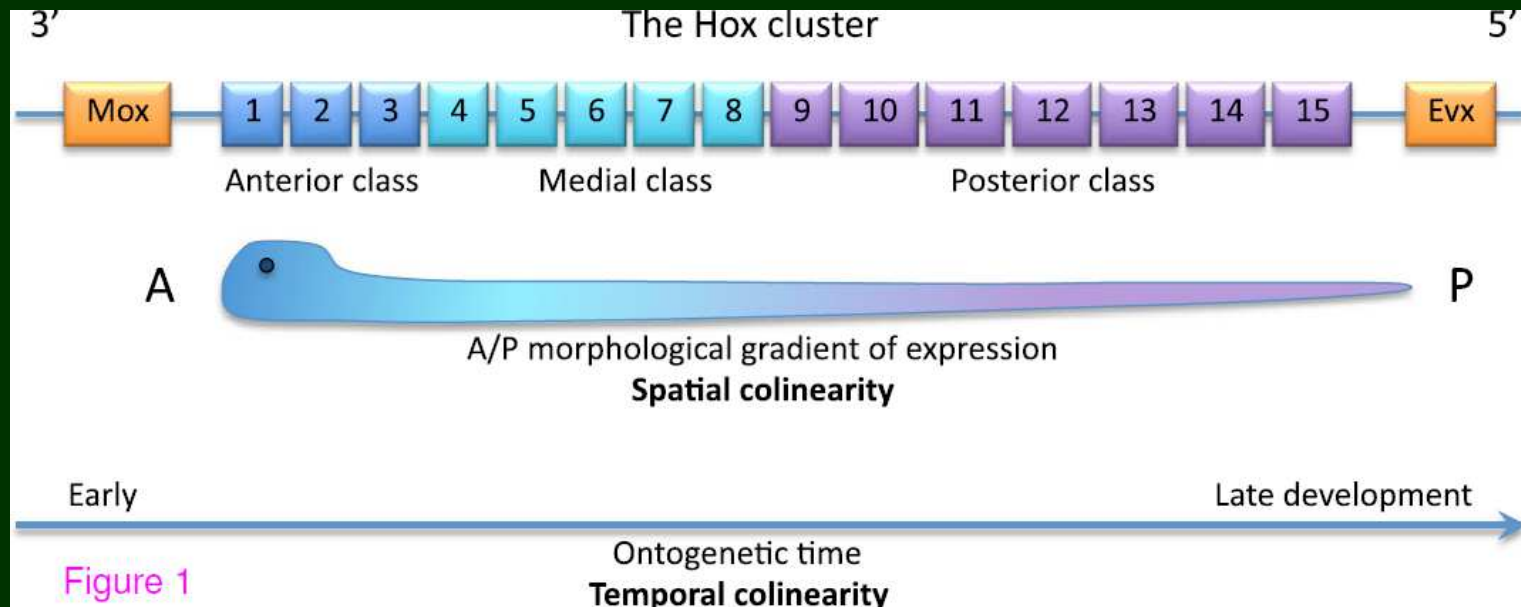


Nervová soustava bilaterií

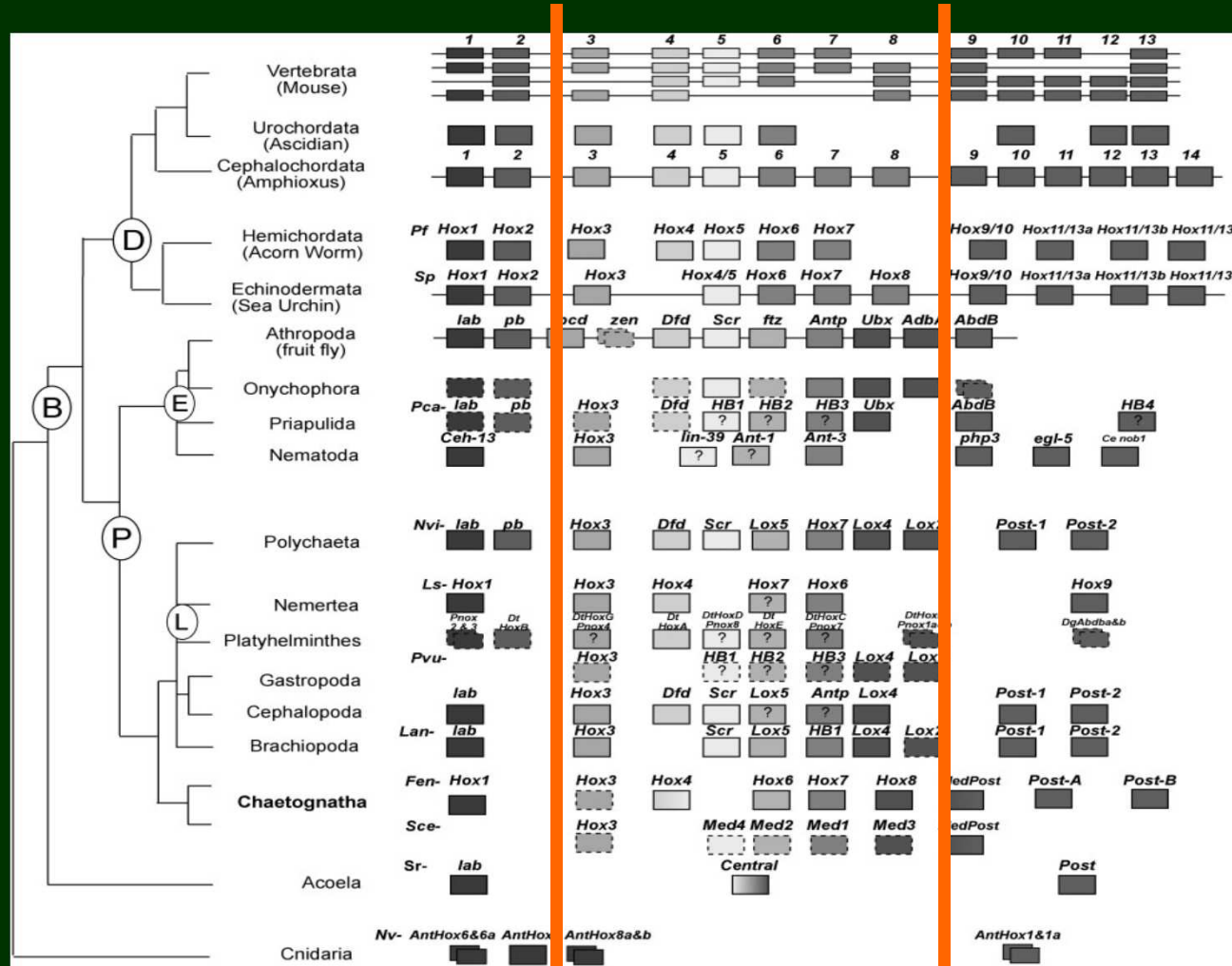


Hox komplexy

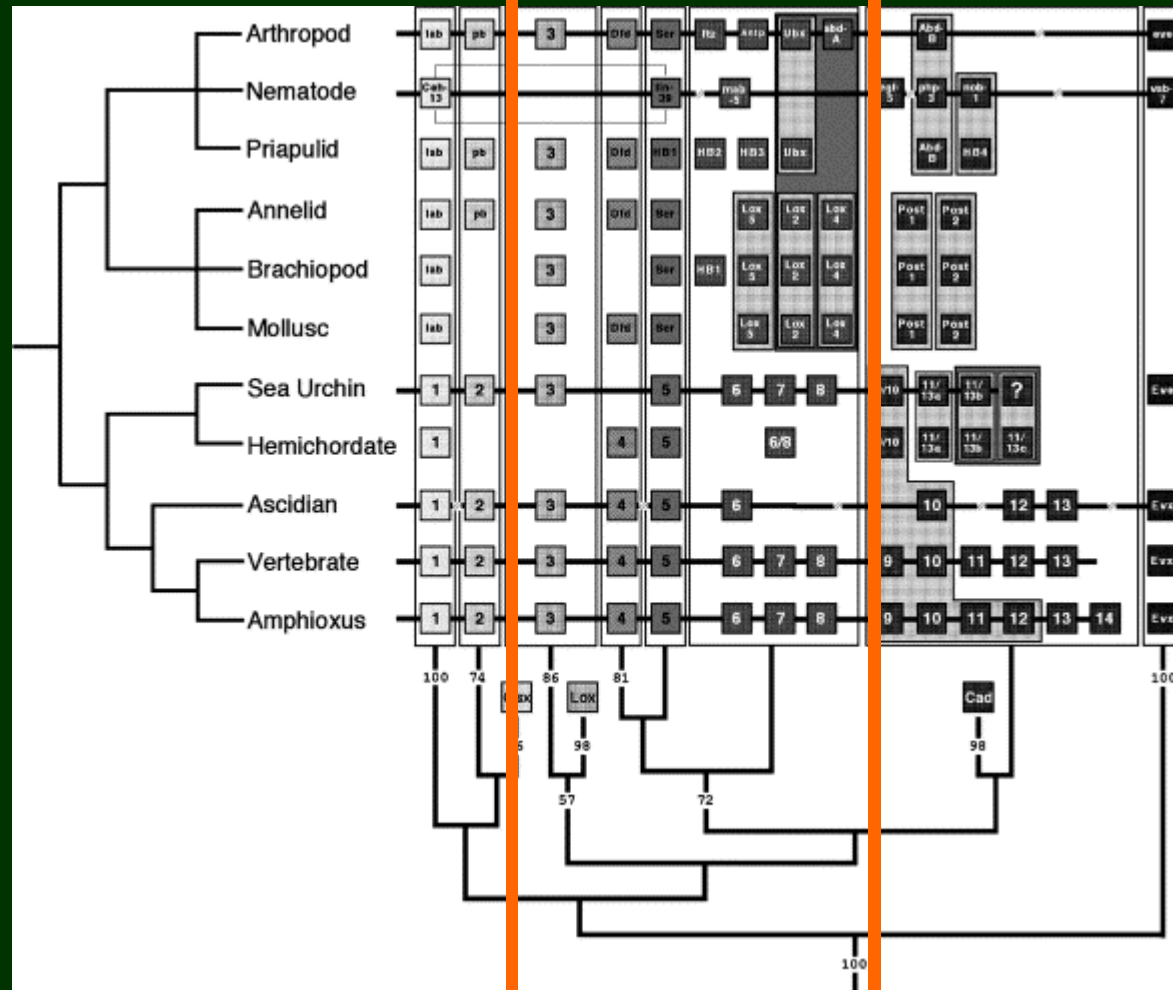
- Hox geny jsou **homologické** nejmíň u všech bilaterií
- tvoří lineární komplexy
- prostorová i časová **kolinearita** (3' = anteriorní a časný)
- “**zootyp**”



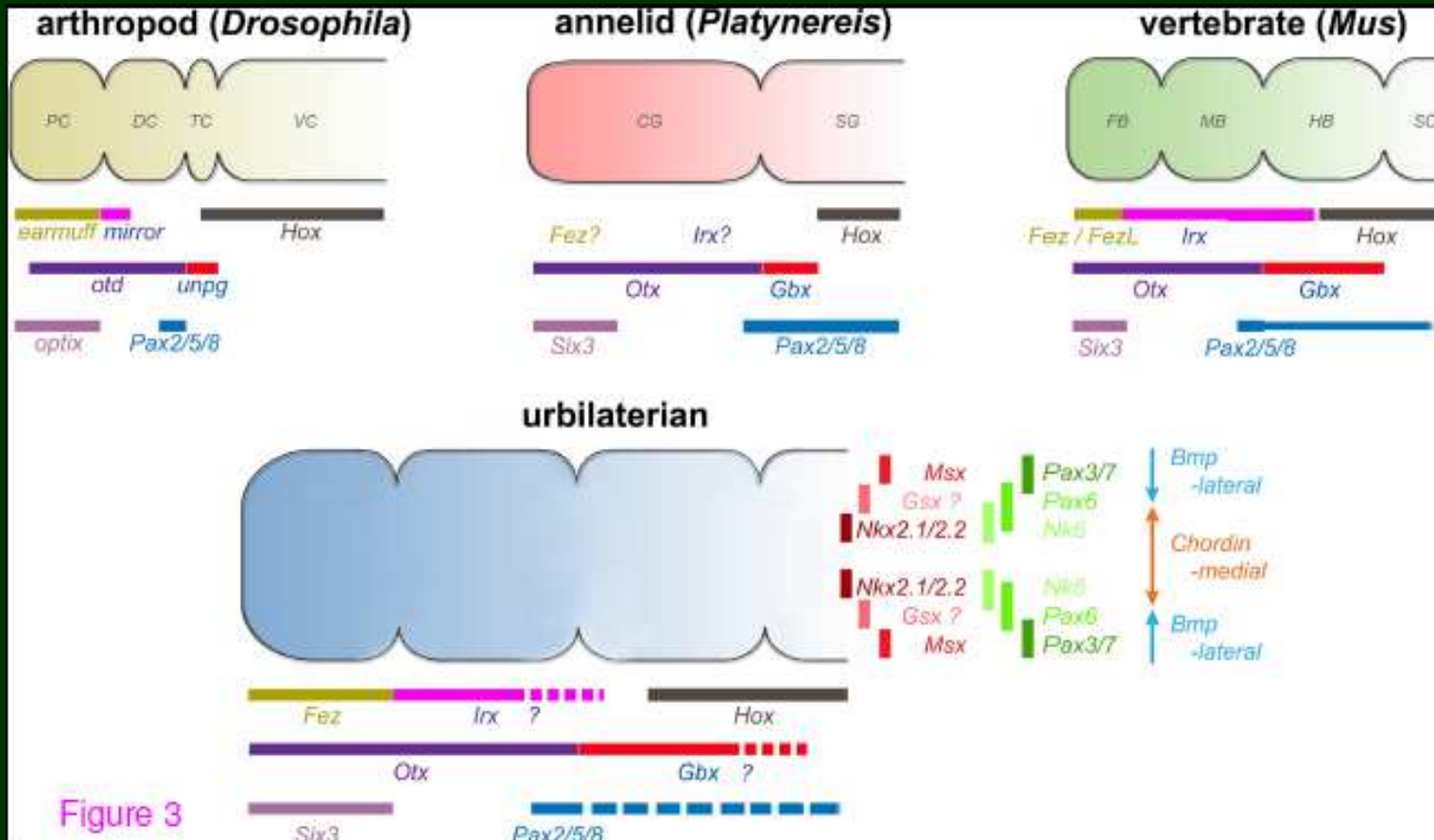
Evoluce Hox komplexů u metazoí



Evolution of Hox genes in two dimensions (orthology x paralogy)



Hox a homologie NS

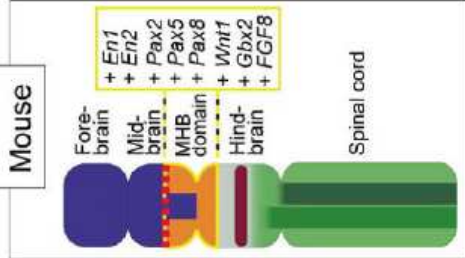


Bilateria

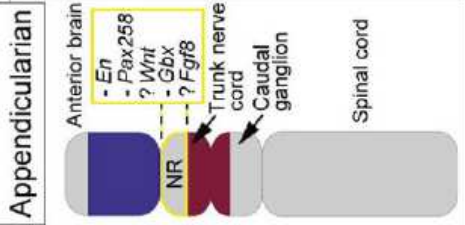
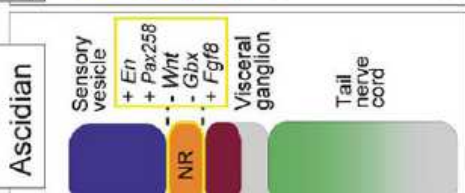
Deuterostomia

Protostomia

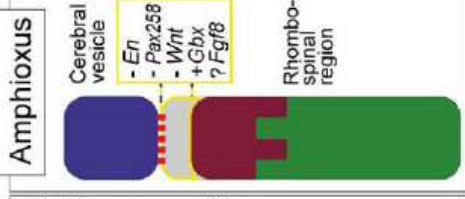
Vertebrata



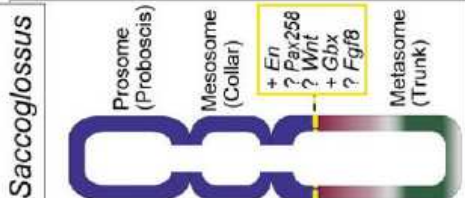
Urochordata



Cephalochordata



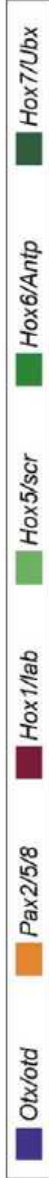
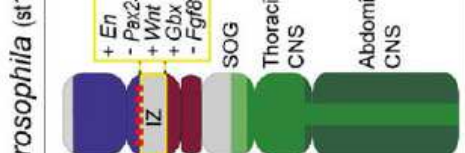
Hemichordata



Annelida

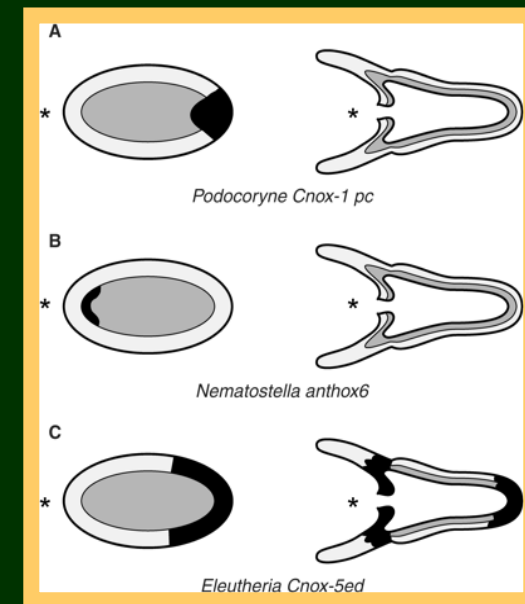
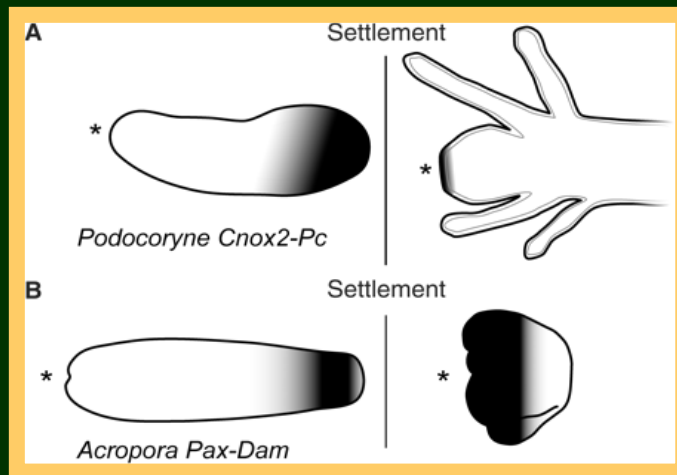


Arthropoda

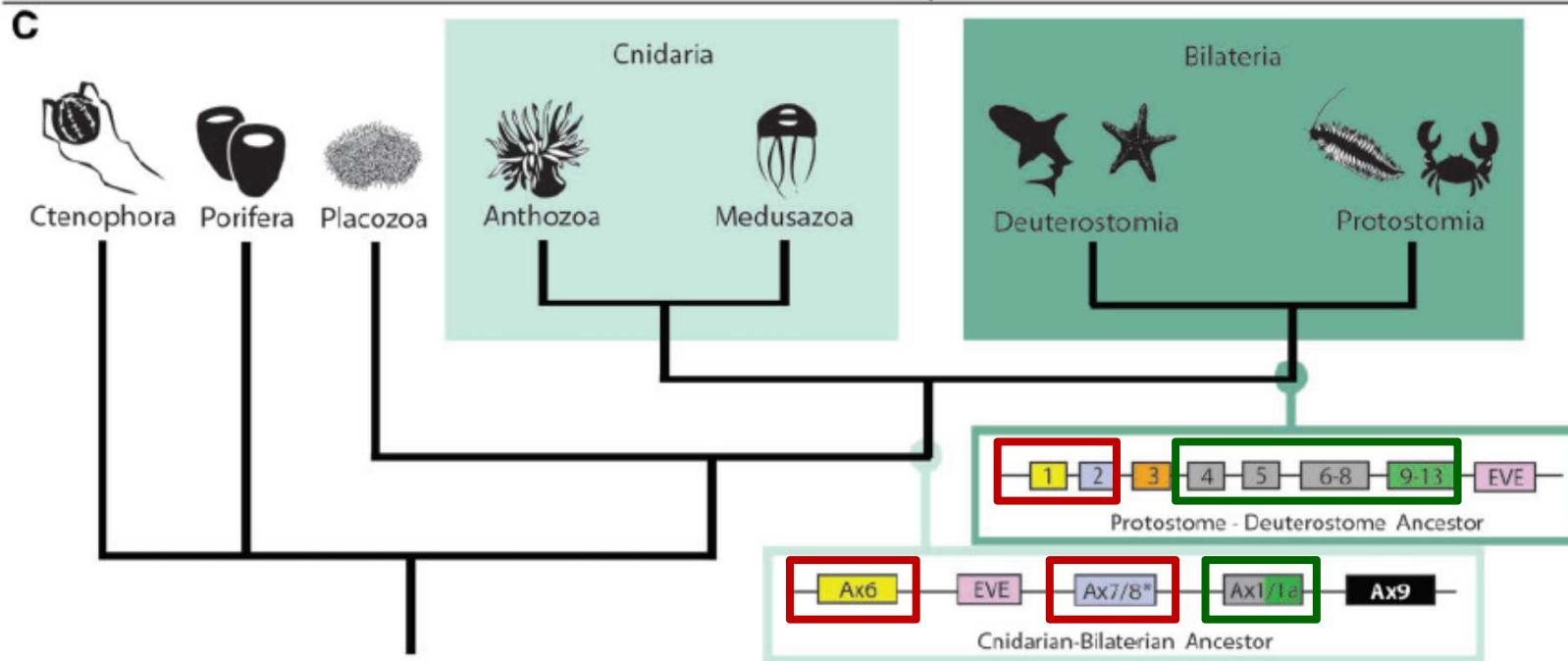
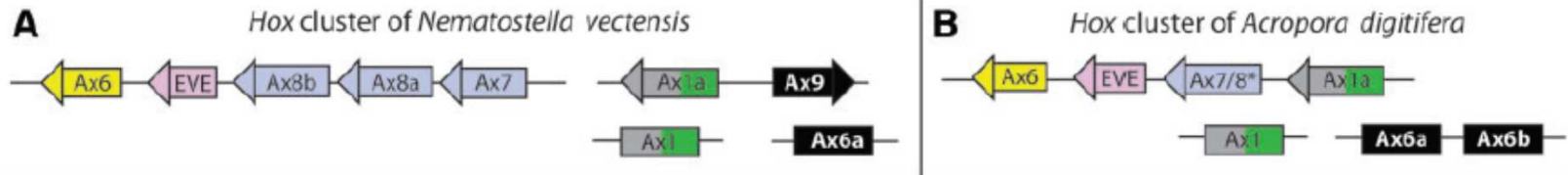


Hox geny u žahavců

- tvoří komplexy (pouze jako výsledky pozdější duplikace)
- některé skupiny Hox genů chybí úplně
- jejich expresní zóny neodpovídají situaci u bilaterií
- jejich expresní zóny vykazují podstatně větší diverzitu než u bilaterií

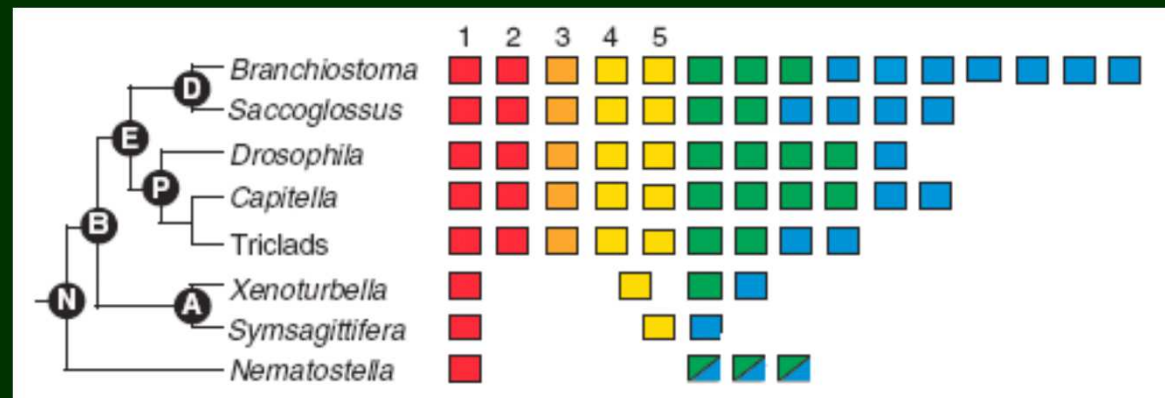
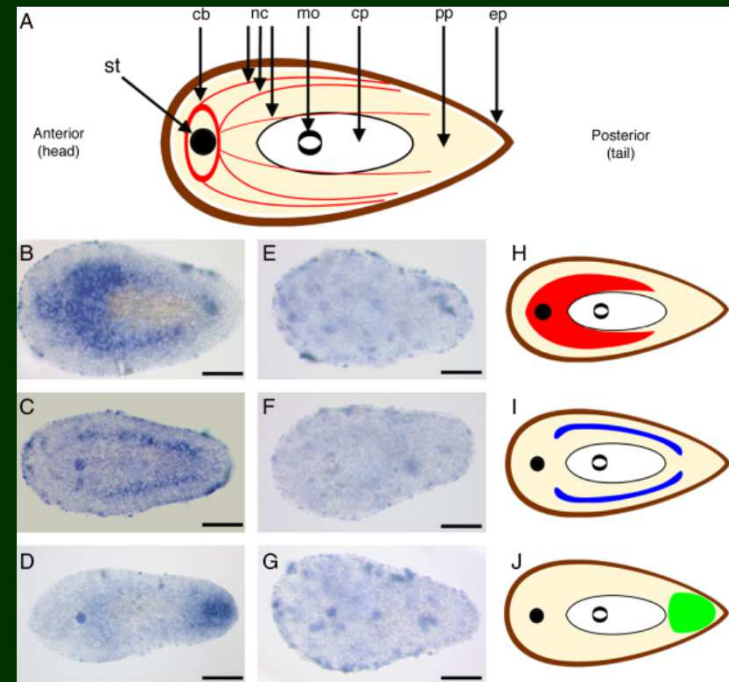


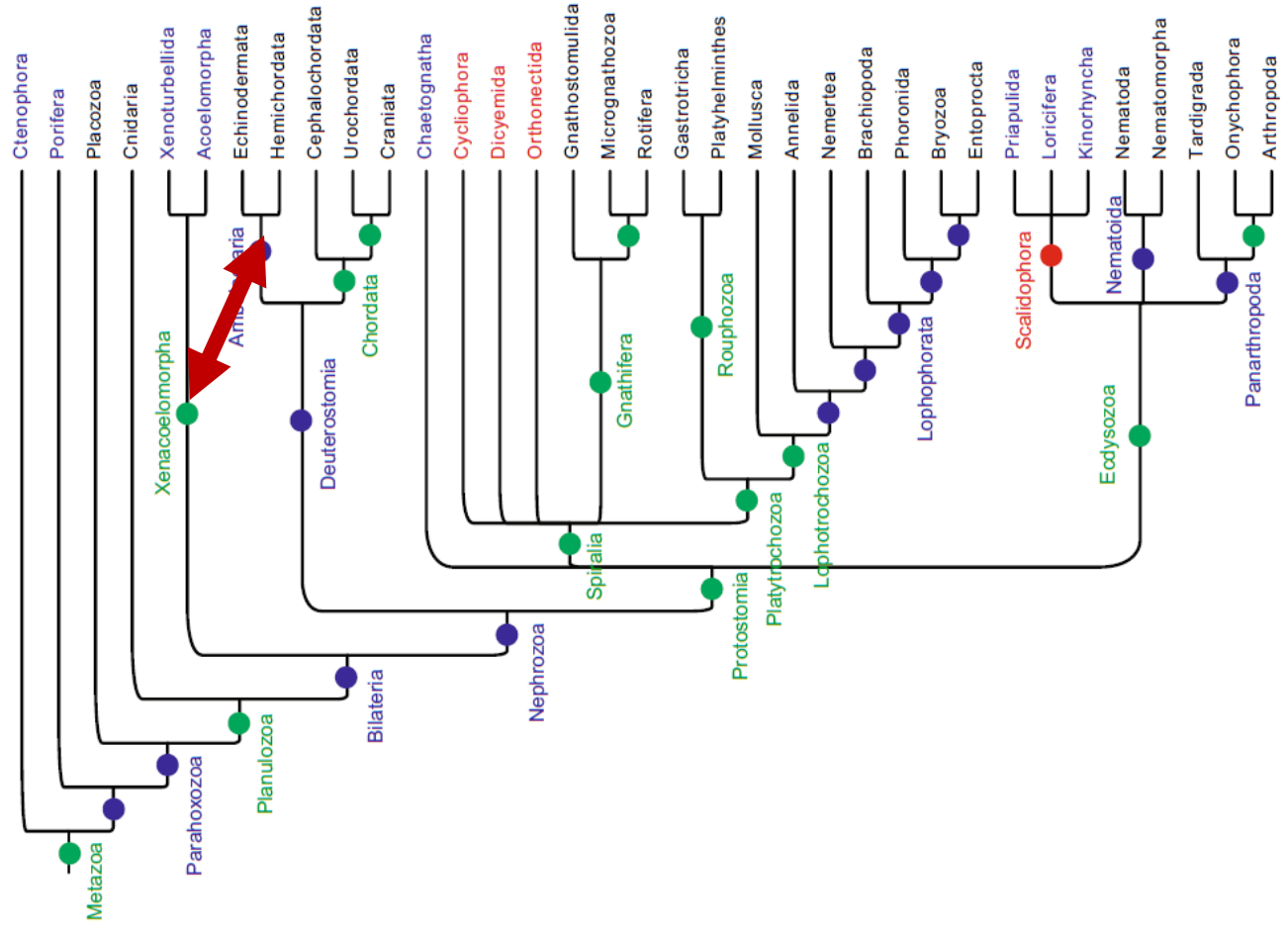
Hox komplex žahavců



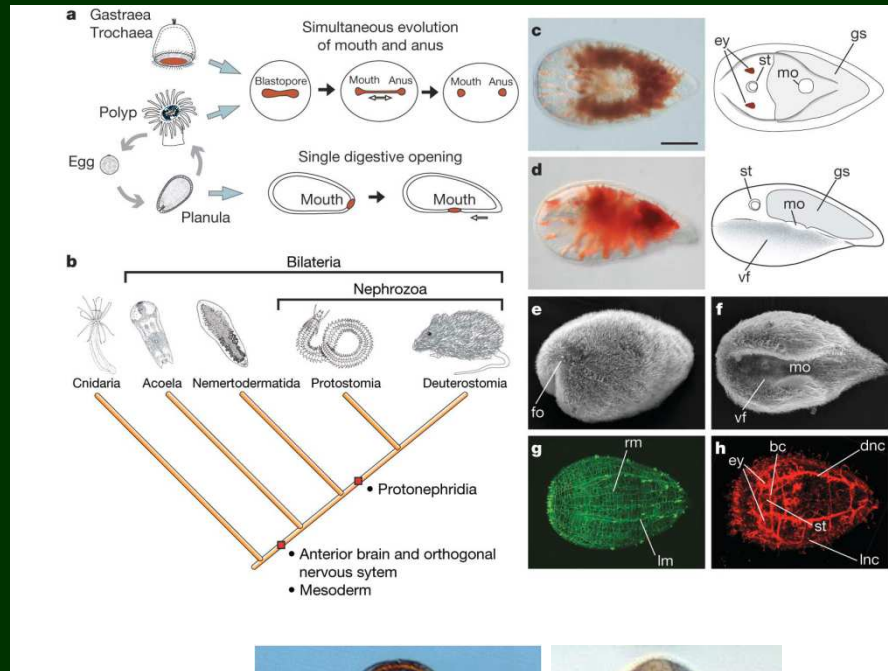
Evoluce Hox systému

- 1. **Cnidaria** ~2-4 geny: A+?C/P (nejasná kolinearita)
- 2. **Acoela** 3 geny: A+C+P
- 3. **Nemertodermatida**: ?+2C+P
- 3. **Xenoturbellida** 4-5 genů: A+2(3)C+P
- (Porifera a Ctenophora 0, Placozoa ~1)





Acoelomorpha



(a) (b)

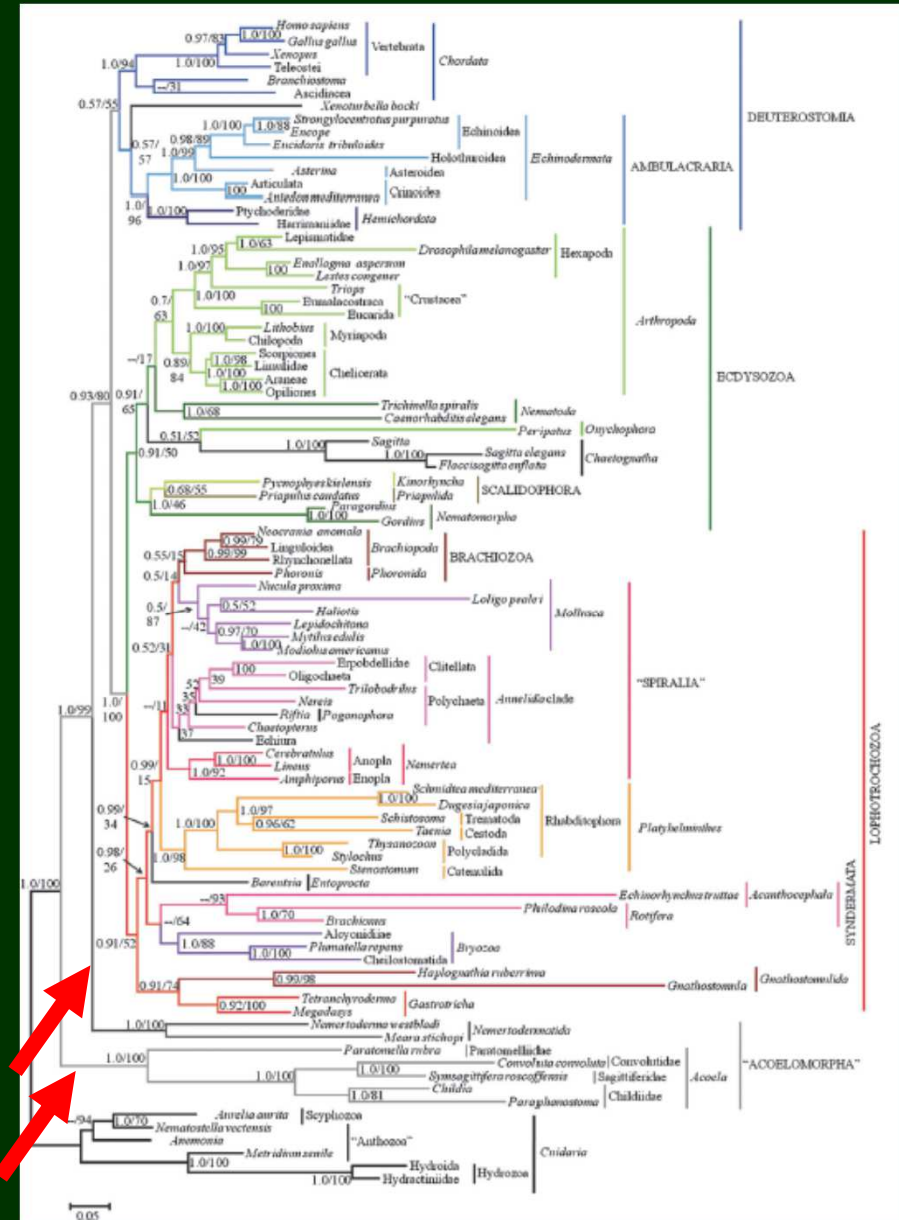
(c)

character	genes	old	new
AP patterning	<i>HOX</i>		
DV patterning	<i>TGFβ/BMP 2/4</i> <i>sog/dpp</i>		
posterior patterning	<i>evx, cdx</i>		
central nervous system	<i>otx, emx, six3/6, HOX1</i>		
photoreception 'eyes'	<i>PAX6, RX, opsin</i>		
heart	<i>tinman</i>		
segmentation, segmentation clock	<i>hairy, engrailed, notch/delta</i>		
regionalized through gut	<i>HNFβ 3, GATA factor</i> <i>goosecoid, brachyury</i>		
appendages	<i>Distal-less/DLX</i>		

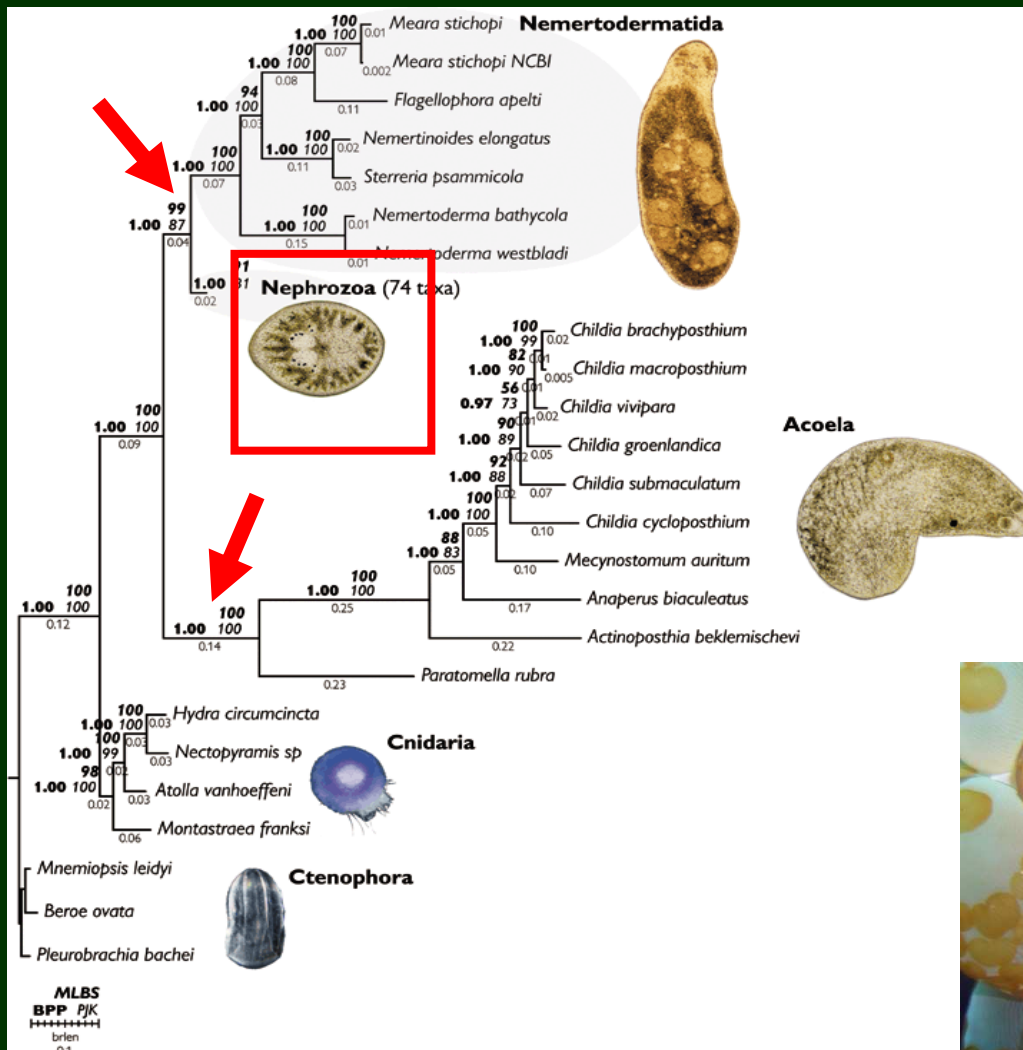
- Acoela + Nemertodermatida

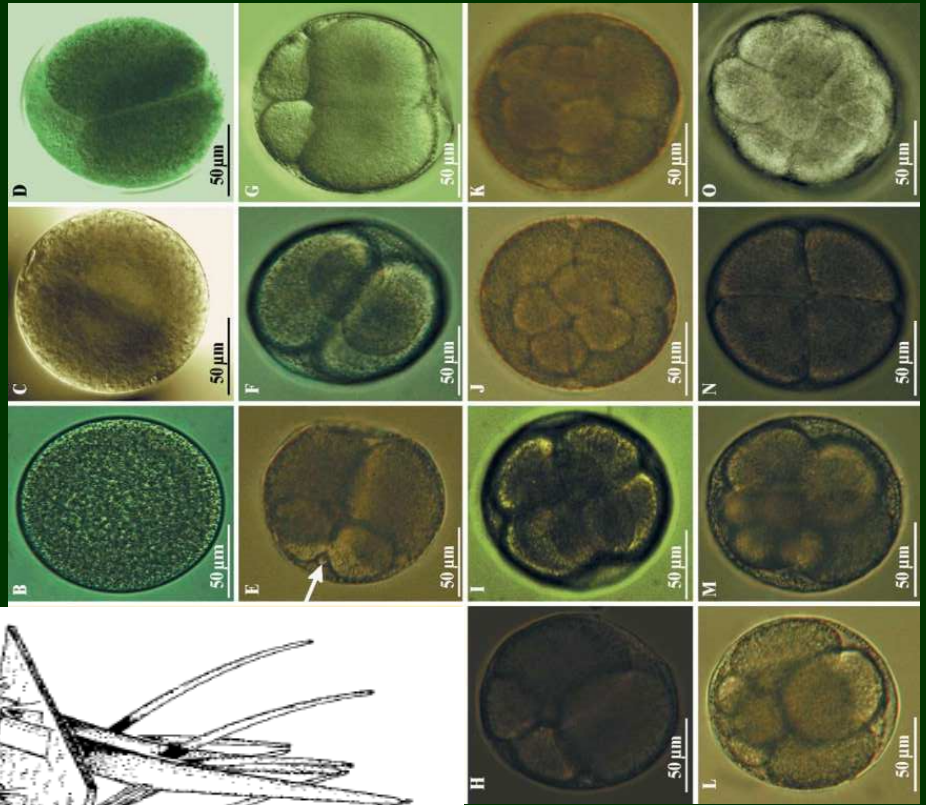
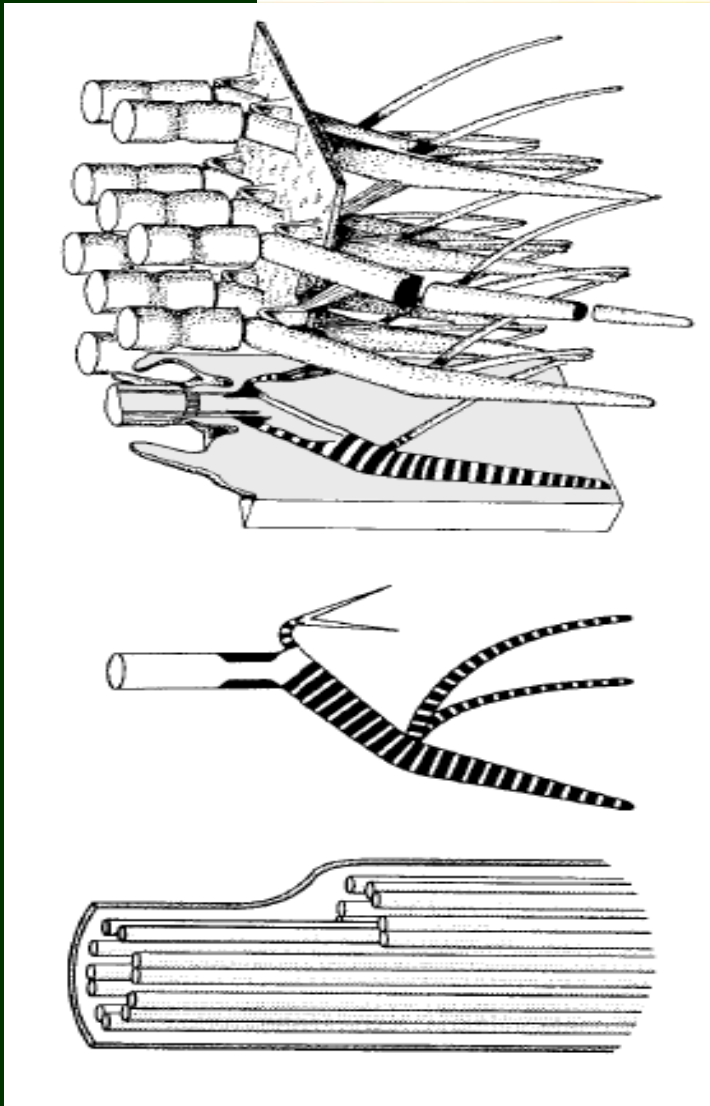
Acoelomorpha: bazální a parafyletická???

13 jaderných
genů



Acoelomorpha: bazální a parafyletická???





Xenacoelomorpha: Nemertodermatida

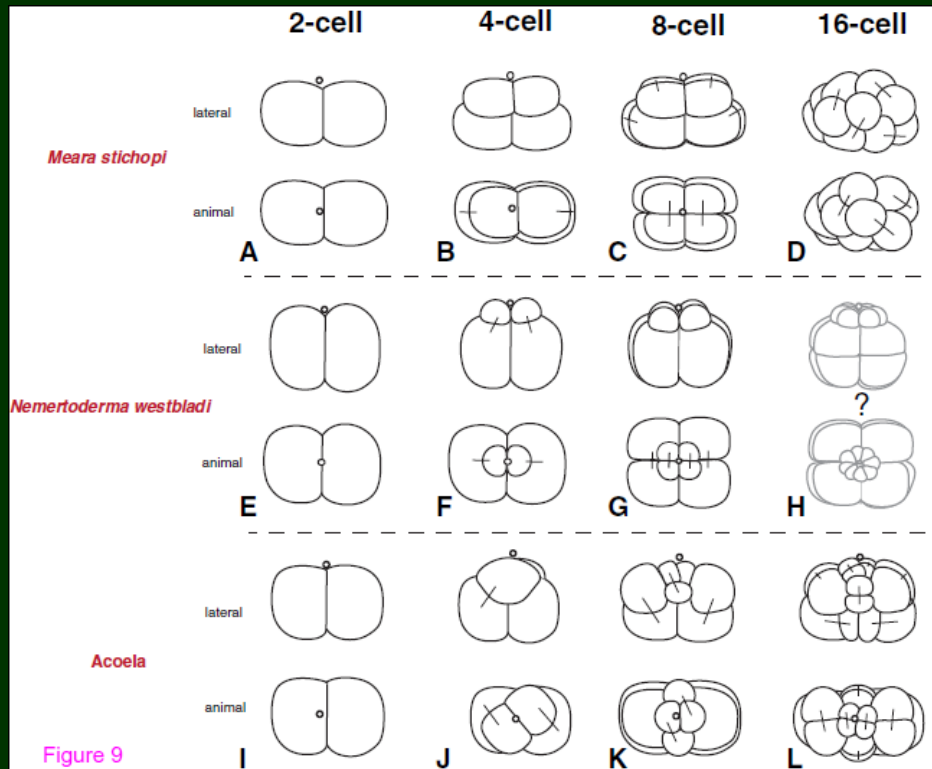
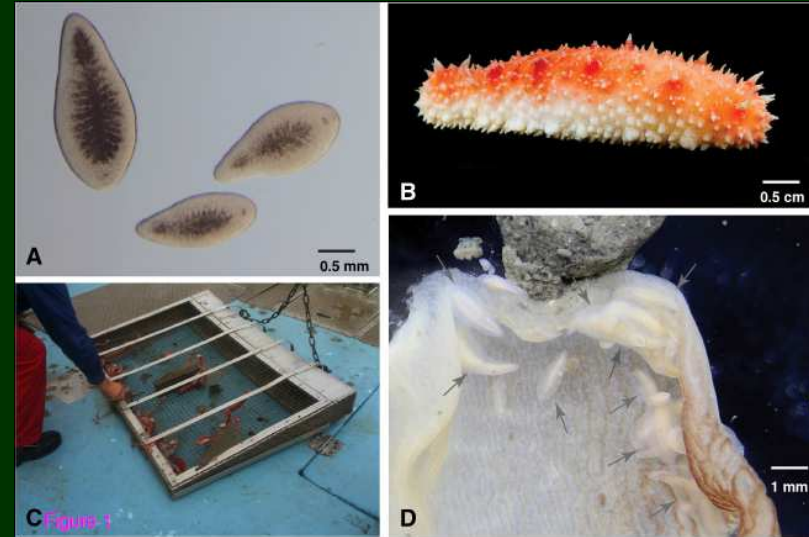
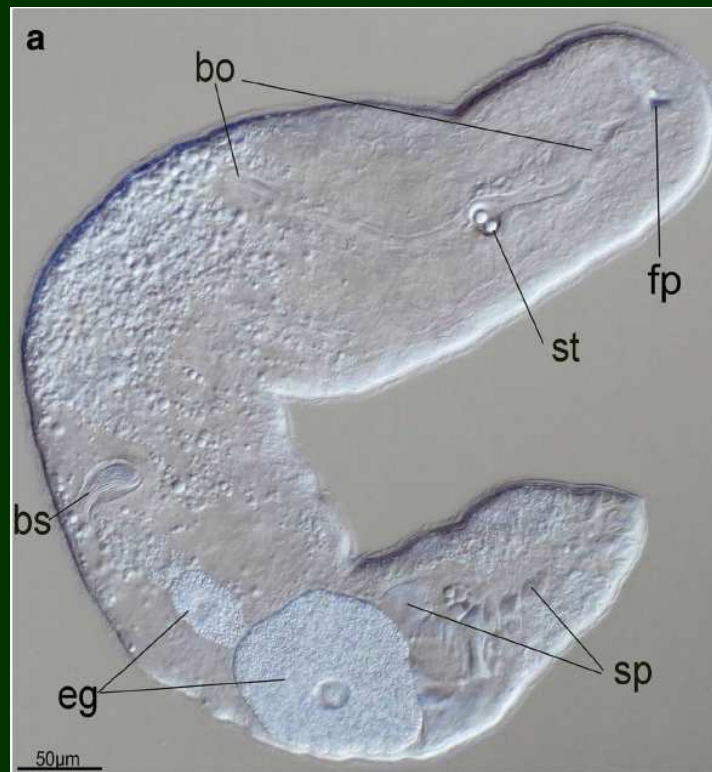


Figure 9

Meara – komezál v přední části TS sumýše
Parastichopus

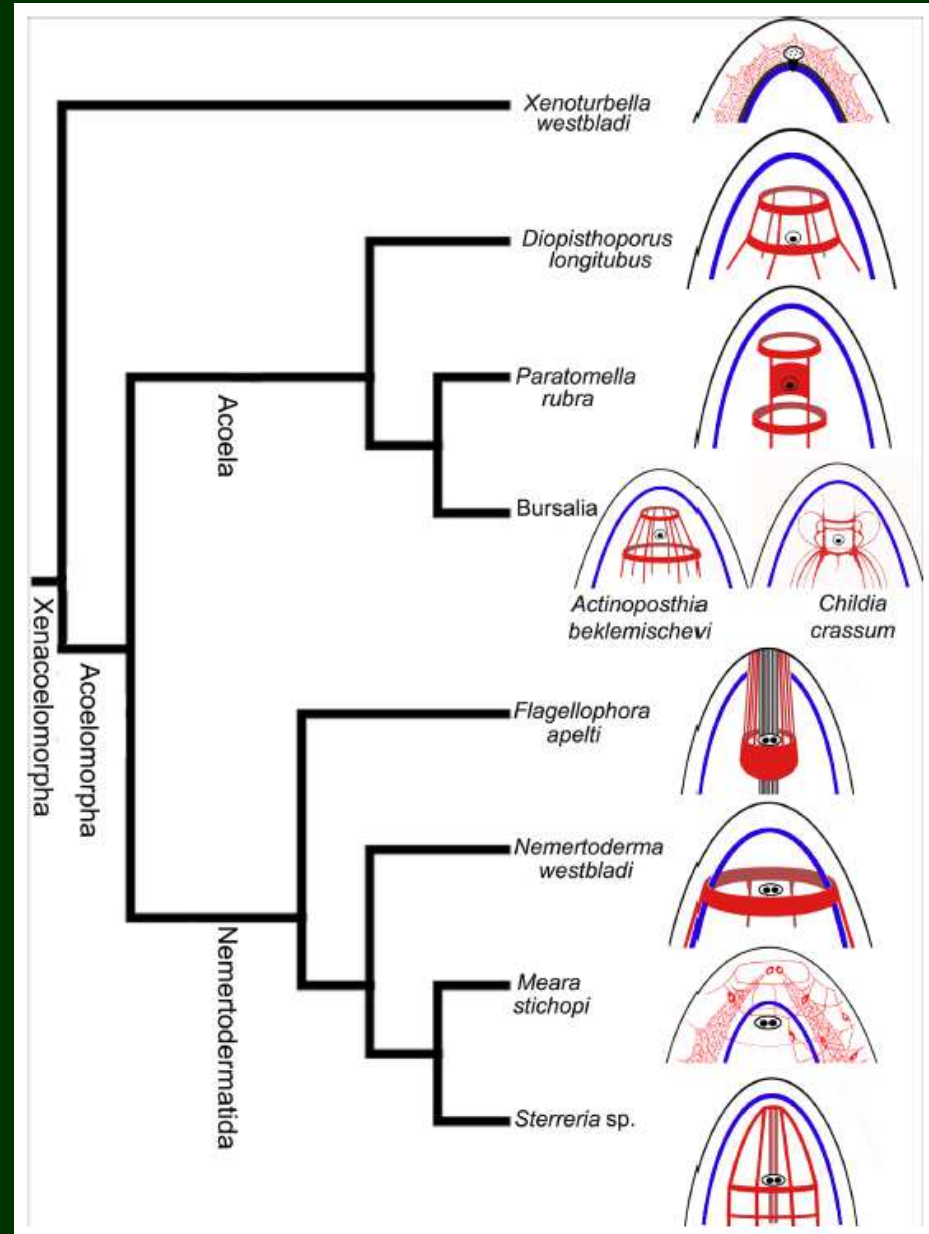
Xenacoelomorpha

- statocysty s pohyblivými buňkami (statokoniemi), statolity ve statokoniích
- statokonie monoflagelátní u X, aflagelátní u Acm (1 u Ac, 2 u Nd)



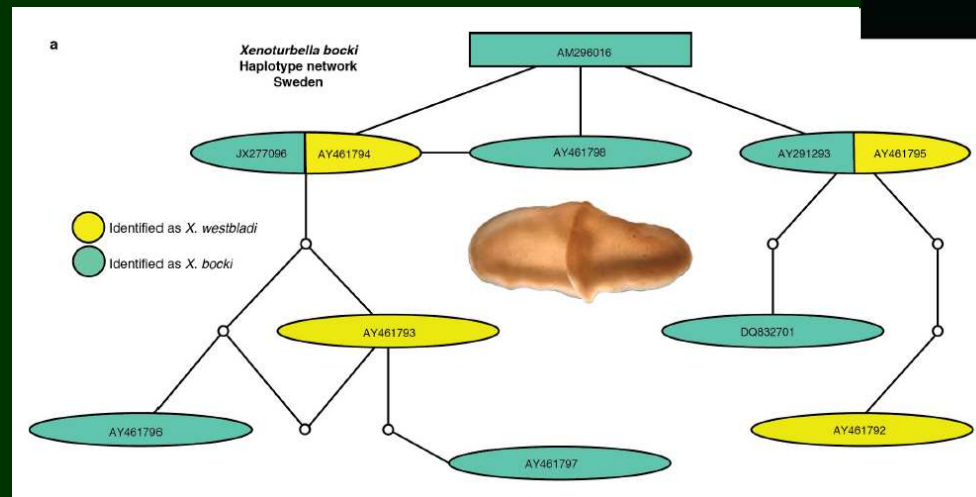
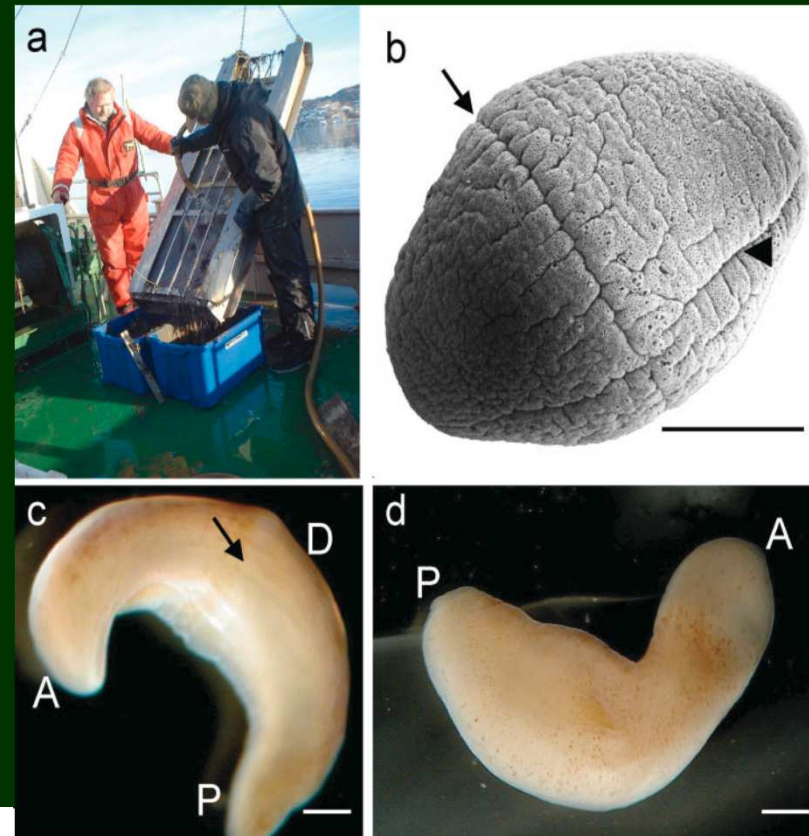
Acoelomorpha NS

- odpovídá předkům s jednoduchým mozkem
- obrovská diverzita

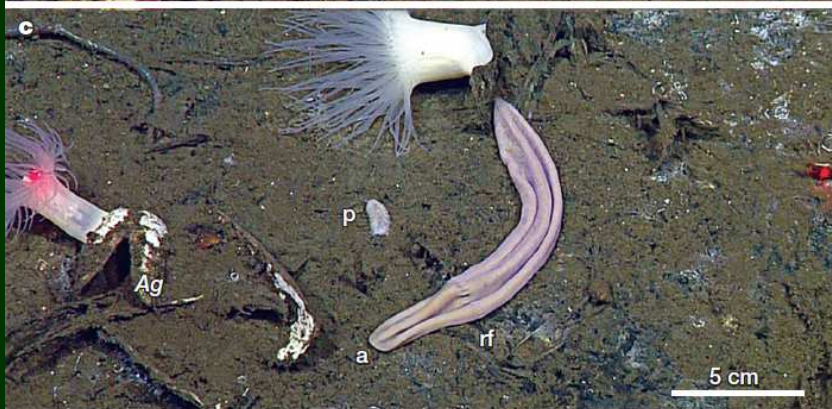


Xenoturbellida

- *Xenoturbella bocki*.
Gullmarsfjord, 50-150 m
- ryje tunely (v laboratoři
hluboké až 15 cm)
- žíví se patrně ranými stadii
mlžů nebo jejich slizem/výkaly

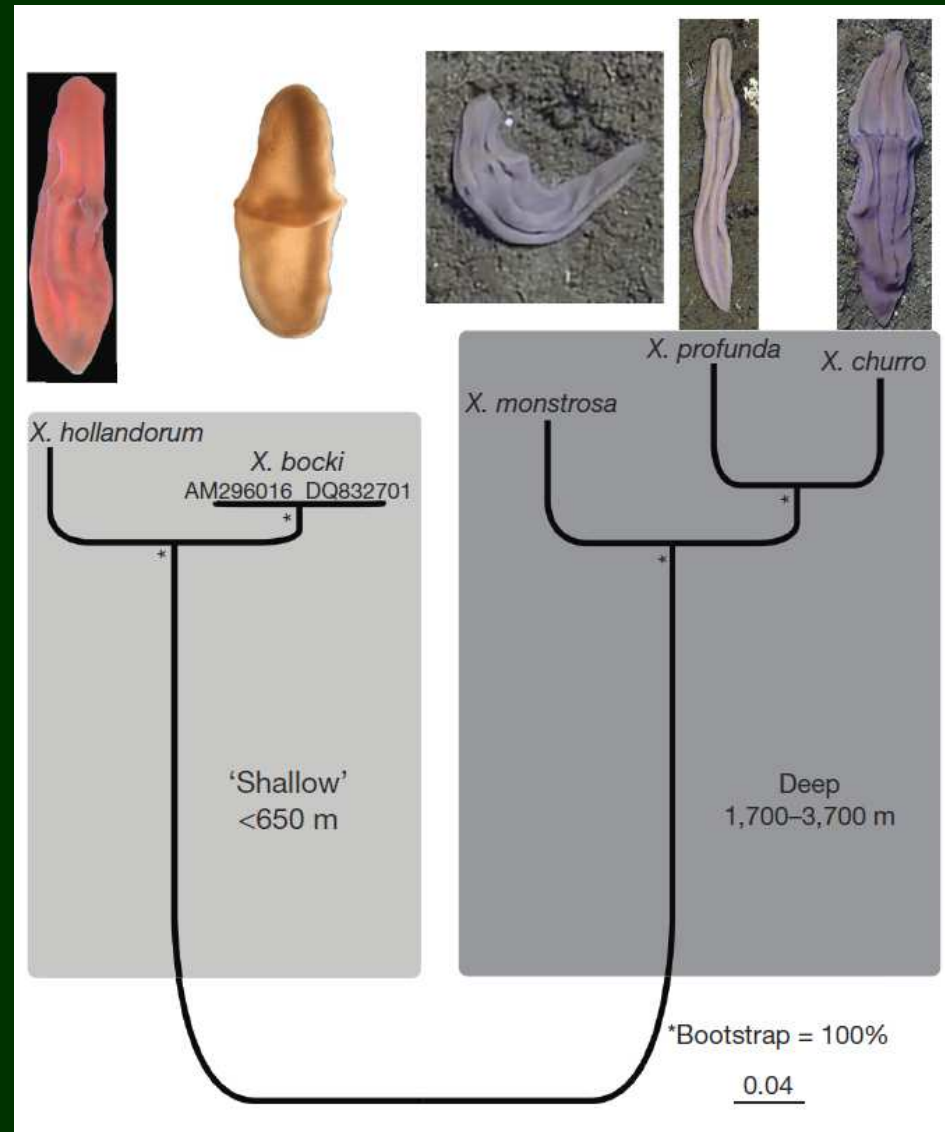


Xenoturbellida



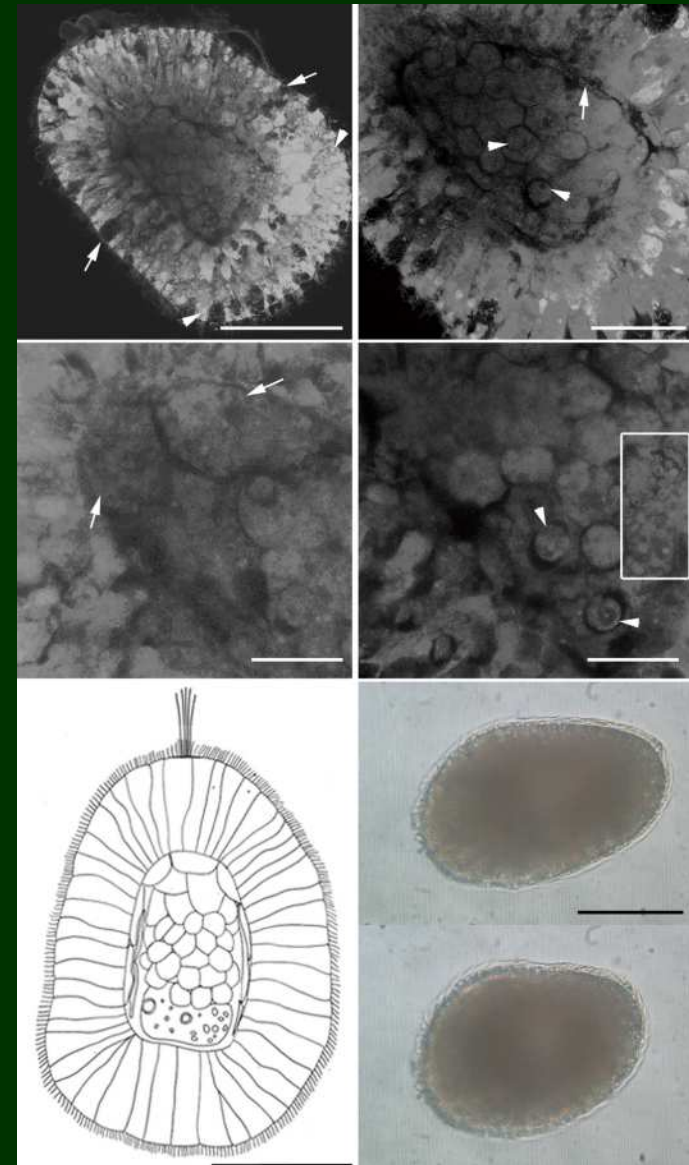
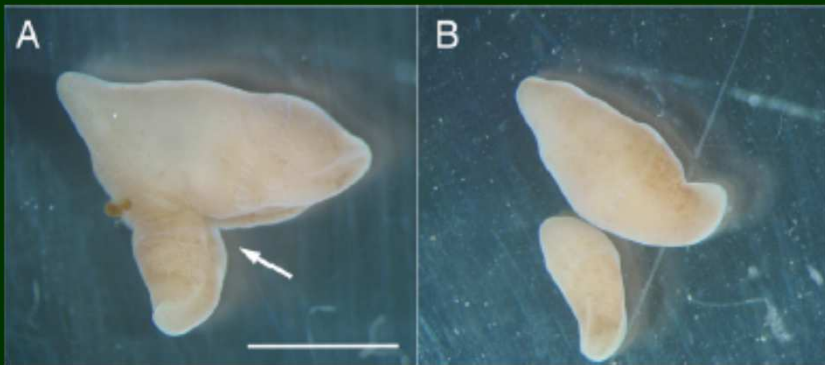
Xenoturbellida

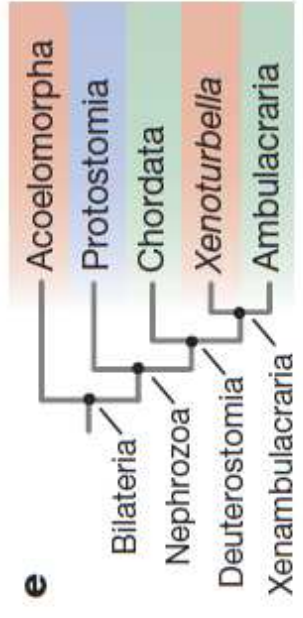
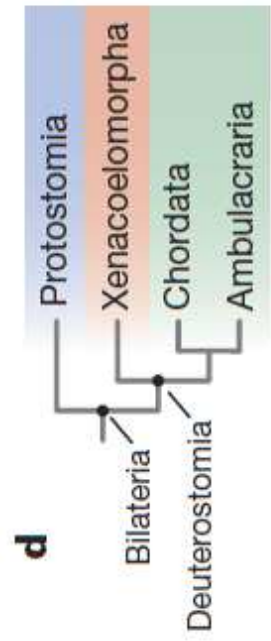
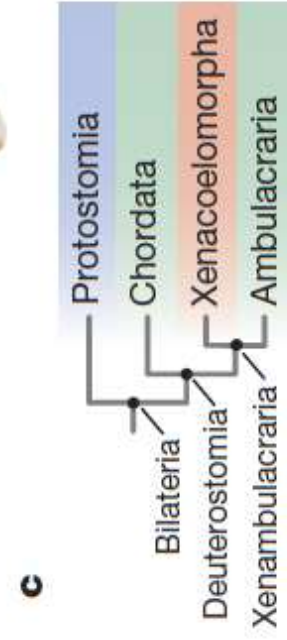
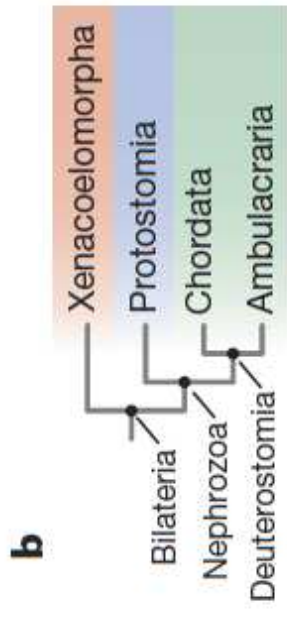
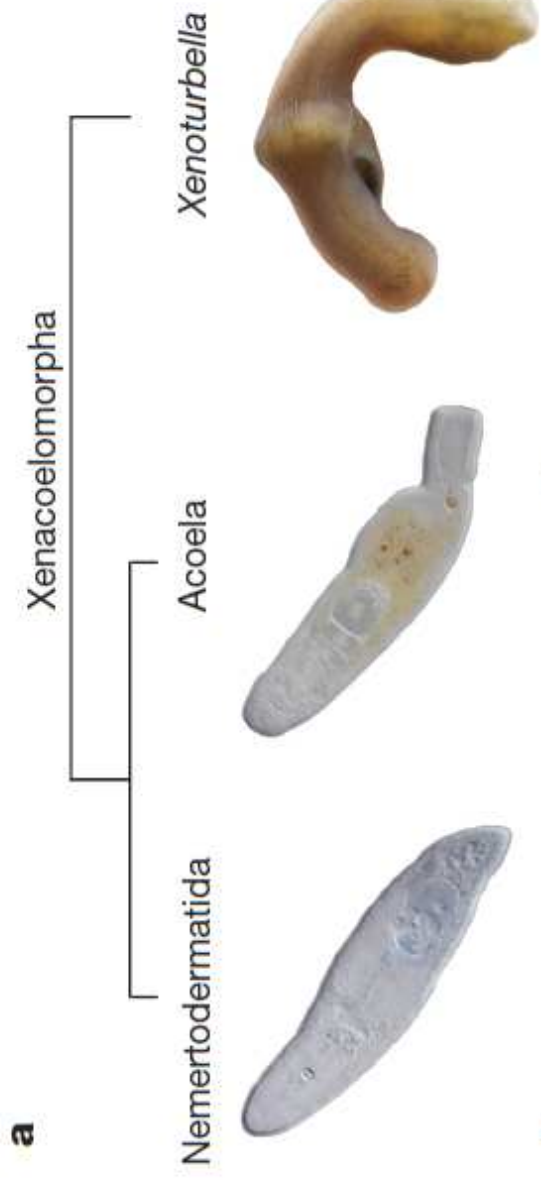
- hlubokomořské formy až 20 cm



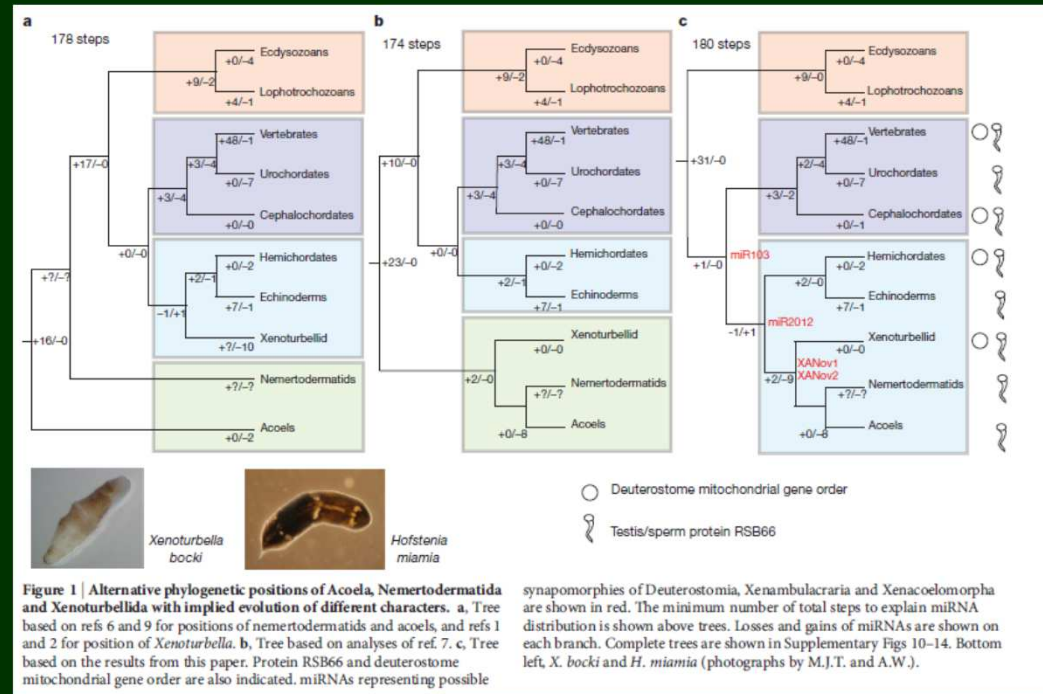
Xenoturbellida

- přímý vývoj – obrvená plovoucí stadia s apikálním orgánem ~ planula (nežerou)
- x druhotně zjednodušené larvy bilaterií obvykle mají zbytky trávicí soustavy (ústa, řiť) či obrvených pásů
- asexuální dělení??? (regenerace – nový jedinec musí obsahovat „statocystu“)





microRNA

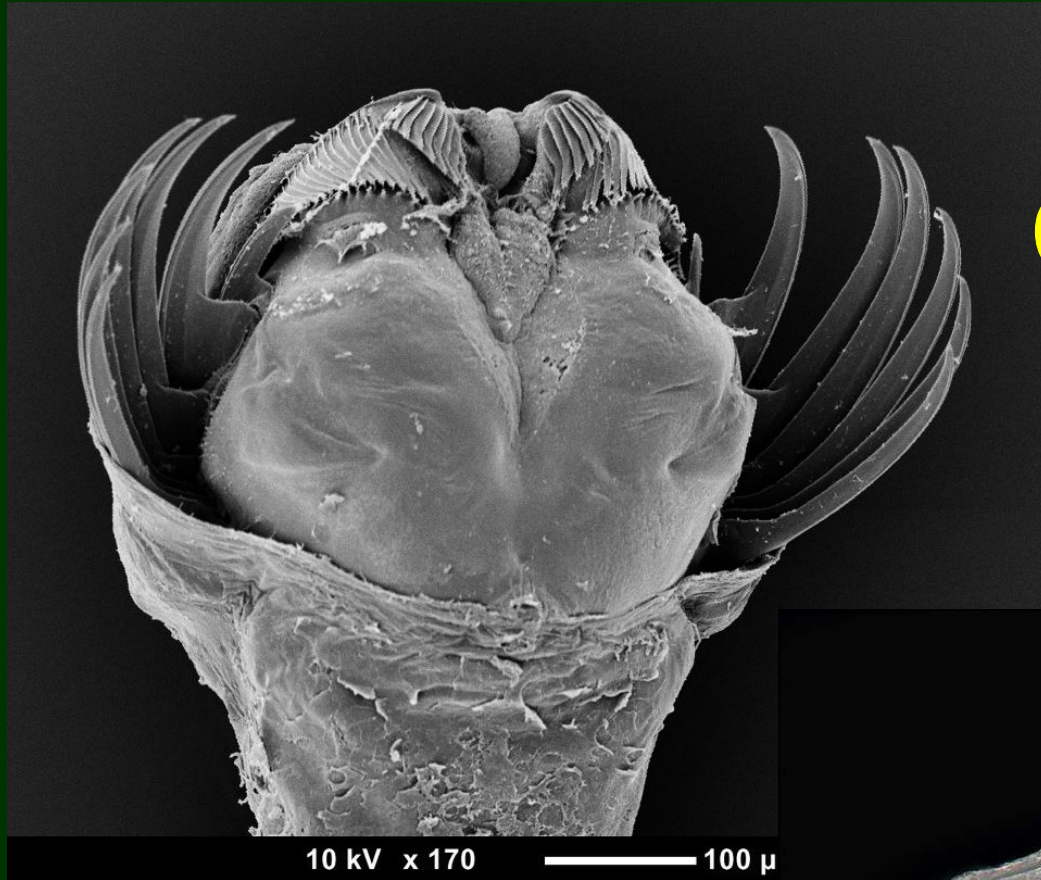


- zásadní rozdíly v prezenci/absenci miRNA i u různých druhů Acoela (*Hofstenia x Symysagittifera*) → velké redukce miRNA
- x specifické deuterostomní miRNA má *Xenoturbella* a někdy i Acoela

Nephrozoa = Eubilateria

- 1. Hyponeuria
 - 1.1. **Protostomia**
 - 1.1. Spiralia
 - 1.2. Ecdysozoa
 - 1.2. Chaetognatha
 - 2. **Deuterostomia**
 - 2.1. Ambulacraria
 - 2.2. Chordata
-
- trávící trubice s řitním otvorem, nefridia, mozek, planktonní larvy ???

Chaetognatha

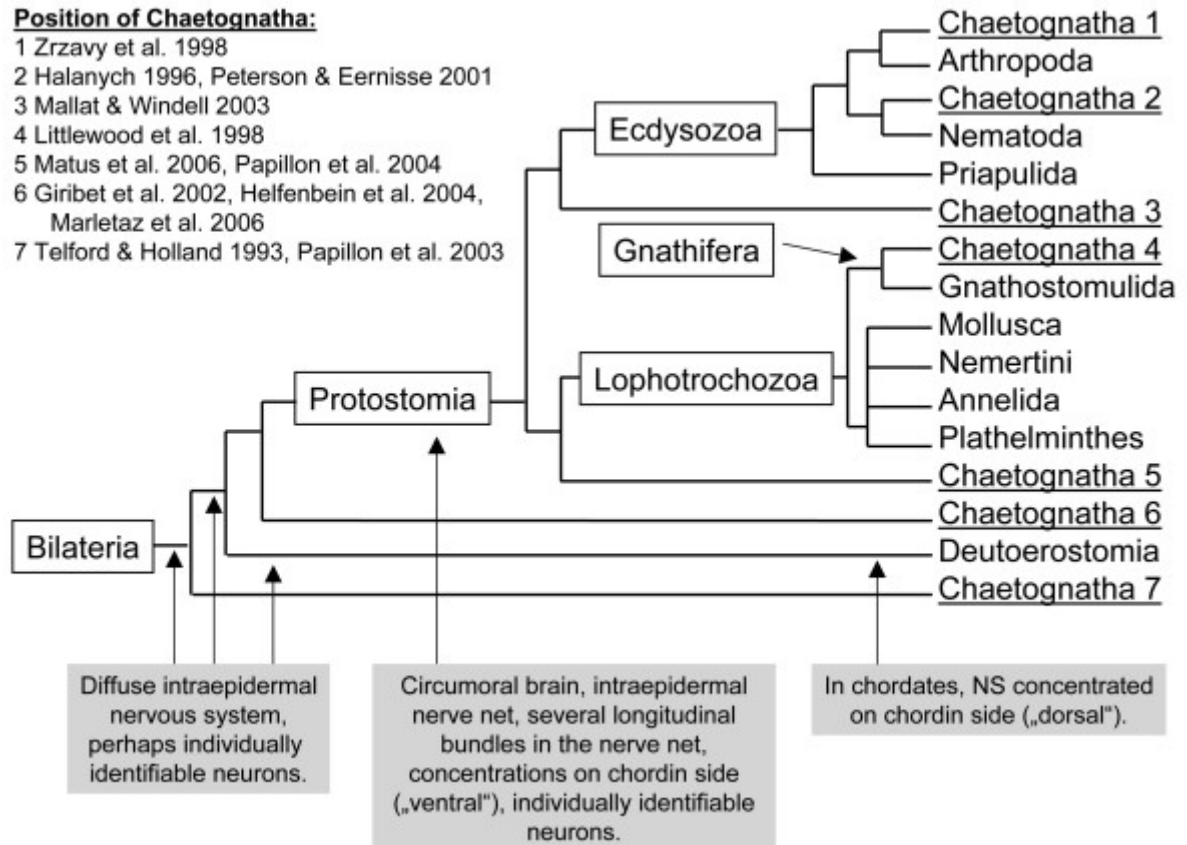


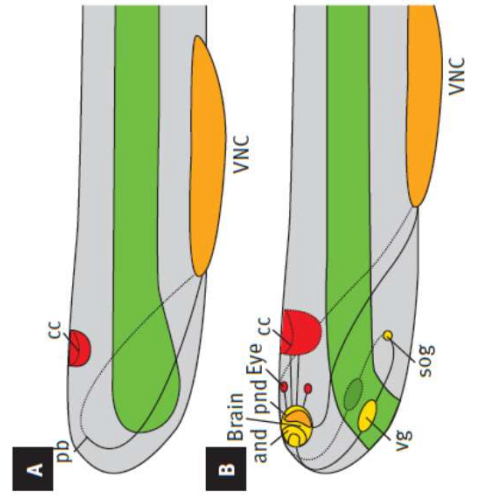
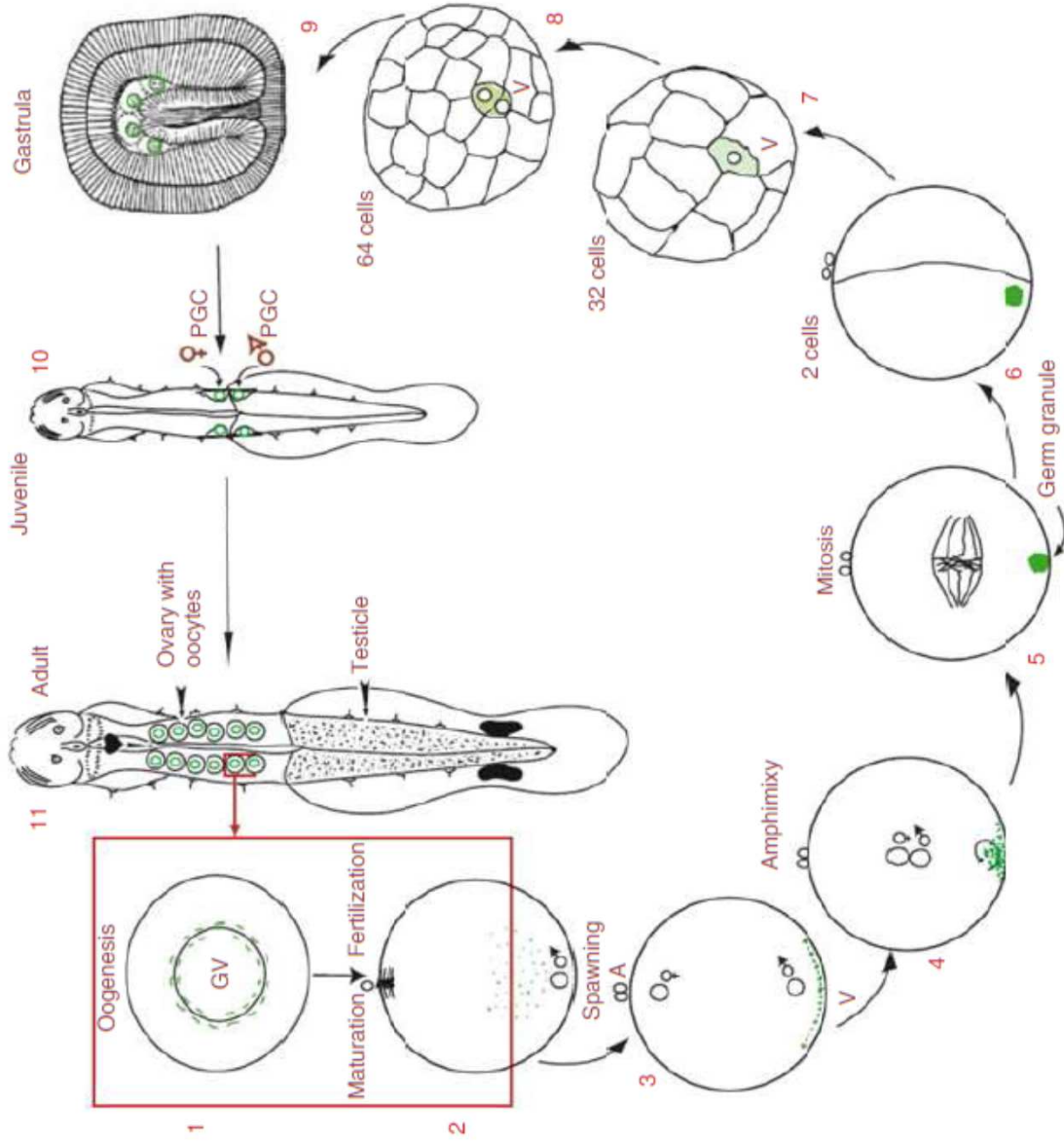
Chaetognatha



Position of Chaetognatha:

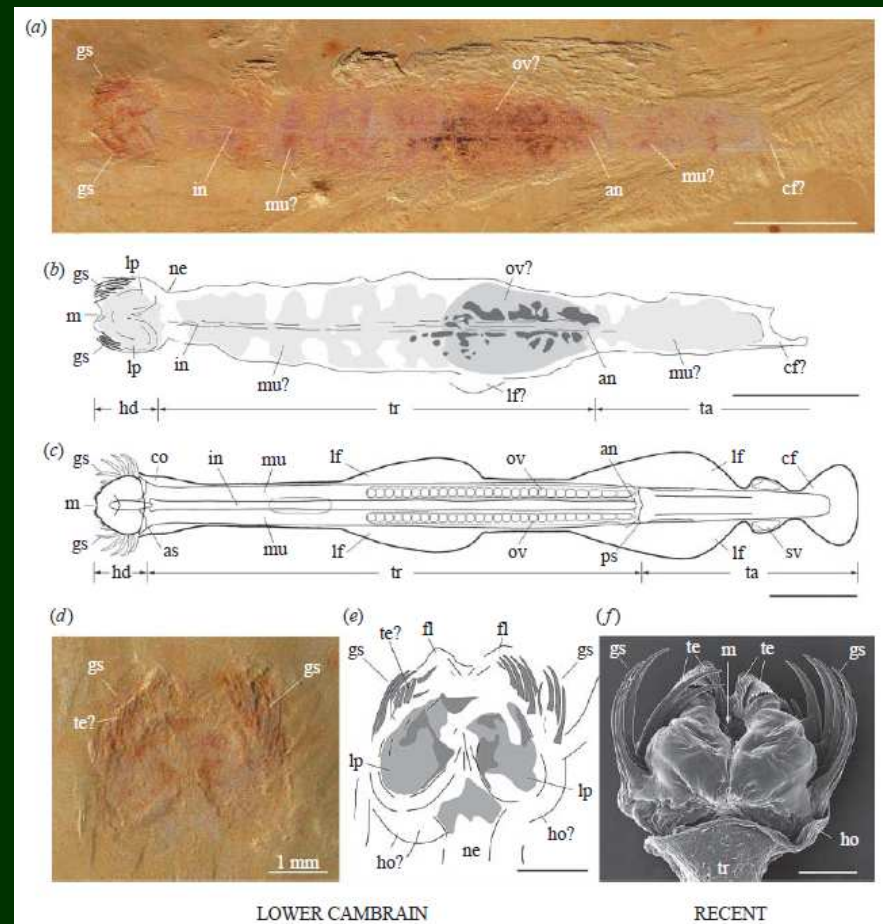
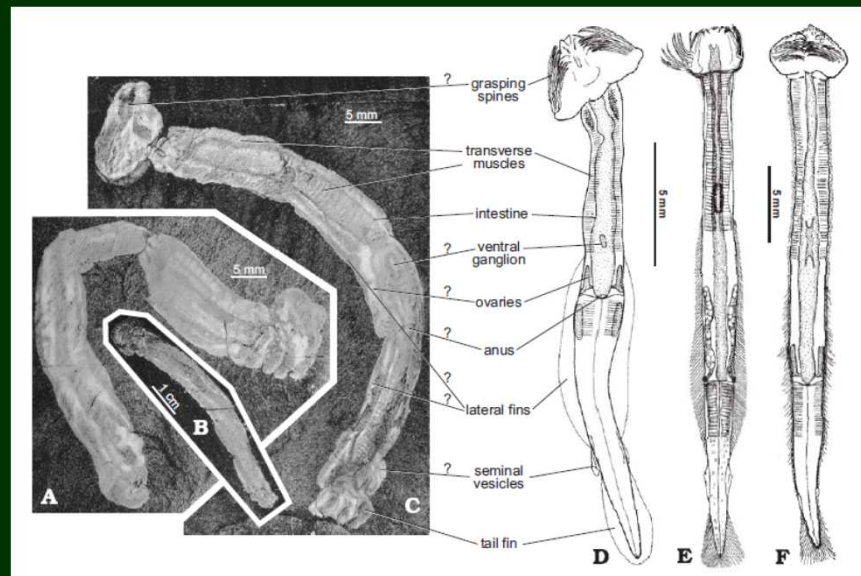
- 1 Zrzavy et al. 1998
- 2 Halanych 1996, Peterson & Eernisse 2001
- 3 Mallat & Windell 2003
- 4 Littlewood et al. 1998
- 5 Matus et al. 2006, Papillon et al. 2004
- 6 Giribet et al. 2002, Helfenbein et al. 2004, Marletaz et al. 2006
- 7 Telford & Holland 1993, Papillon et al. 2003





Chaetognatha

- fosilní záznam patrně už z kambria (*Protosagitta*, *Oesia*?)
- *Paucijaculum* (karbon)
- (protokonodonty?)
- (*Amiskwia*???)



Chaetognatha

mtDNA

Sagitta crassa, *Zonosa agilis* and *Decaploagitta decipiens* (Chaetognatha: Aphragmophora: Sagittidae):

DRNA	cox2	cox3	nad1	nad3	nad2	cox3	srRNA	nad4	nad6	nad5	nad4L	cob
------	------	------	------	------	------	------	-------	------	------	------	-------	-----

Sagitta orfata (Chaetognatha: Aphragmophora: Sagittidae):

DRNA	cox2	cox3	nad1	nad3	nad2	cox3	srRNA	nad4	nad5	nad4L	cob	nad6
------	------	------	------	------	------	------	-------	------	------	-------	-----	------

Parapadilla gotoi (Chaetognatha: Phragmophora: Spadellidae):

DRNA	cox2	cox3	nad1	nad3	nad2	cox3	srRNA	nad5	nad4L	nad4	cob	nad6
------	------	------	------	------	------	------	-------	------	-------	------	-----	------

Spadella cephalopora (Chaetognatha: Phragmophora: Spadellidae):

DRNA	cox2	nad1	nad3	nad2	cox3	srRNA	cox1	nad4L	nad4	nad5	cob	nad6
------	------	------	------	------	------	-------	------	-------	------	------	-----	------

Lumbricus terrestris, *Haymeria danovii*, *Whitmania pigra*, *Clymenella torquata*, *Orbinia latreilli*, *Parionys excavatus*, *Nephtys* sp. "San Juan Island" YV-2008, *Pista cruxata*, *Terebellides strocmii*, *Perinereis nuntia*, *Marphysa sanguinea*, *Hirudo nipponia*, *Poecilobdella manilensis*, *Whitmania laevis*, *Eteobdella octocolorata*, *Tonno colex* *birmanicus* and *Metaphire vulgaris* (Annelida):

DRNA	nad1	nad3	nad2	cox1	cox2	atp8	cox3	nad6	cob	atp6	nad5	nad4L	nad4	srRNA
------	------	------	------	------	------	------	------	------	-----	------	------	-------	------	-------

Urechis caupo and *U. unicinctus* (Echidaria); All studied echidarians:

DRNA	cox2	nad6	cob	atp6	nad5	cox1	cox2	atp8	nad4L	nad4	nad2	nad1	nad3	srRNA
------	------	------	-----	------	------	------	------	------	-------	------	------	------	------	-------

Bugula neritima (Bryozoa: Cheilostomatida):

DRNA	nad6	cox1	atp8	atp6	nad1	nad2	cox2	cob	nad4L	nad4	nad5	nad1	cox1	srRNA
------	------	------	------	------	------	------	------	-----	-------	------	------	------	------	-------

Flustrilla dora *hispida* (Bryozoa: Ctenostomata):

DRNA	cox2	nad1	srRNA	nad4L	atp6	cox1	atp8	nad3	nad2	cob	cox3	nad5	nad1	nad6
------	------	------	-------	-------	------	------	------	------	------	-----	------	------	------	------

Phoronopsis harmeri (Brachiopoda: Phoroniformea):

DRNA	srRNA	cox1	atp6	nad3	nad2	cox1	cox2	atp8	nad5	nad4	nad4L	cob	nad6	nad1
------	-------	------	------	------	------	------	------	------	------	------	-------	-----	------	------

Terebratalia transversa (Brachiopoda: Rhynchonelliformea):

DRNA	nad1	nad4	nad4L	cox2	nad3	nad2	cox1	cob	atp8	cox3	atp6	nad5	srRNA	nad6
------	------	------	-------	------	------	------	------	-----	------	------	------	------	-------	------

Terebratalia retusa (Brachiopoda: Rhynchonelliformea):

DRNA	nad1	nad6	cob	nad4L	nad4	nad5	cox3	nad1	nad2	cox1	cox2	atp8	atp6	srRNA
------	------	------	-----	-------	------	------	------	------	------	------	------	------	------	-------

Trichinella spiralis, *Trichuris trichiura*, *Trichuris suis*, *Trichuris discolax* and *Trichuris ovis* (Nematoda):

DRNA	atp6	cox3	atp8	nad3	cox1	nad1	nad2	nad5	nad4	nad4L	nad6	cob	srRNA
------	------	------	------	------	------	------	------	------	------	-------	------	-----	-------

Papilio amabilis, *Papilio xanthus* and *Papilio indolatus* (Mollusca):

DRNA	atp6	nad3	nad5	atp8	nad4	nad6	cox3	srRNA	cox1	nad1	nad2	nad4L	cox2	cob
------	------	------	------	------	------	------	------	-------	------	------	------	-------	------	-----

Oscarella carmelis, *Oscarella microlobata*, *Oscarella viridis*, *Oscarella lobulata*, *Oscarella tuberculata* and *Pseudocorticium jarvisi* (Porifera):

DRNA	cox2	cox1	nad1	atp8	atp6	cox3	srRNA	nad8	nad6	nad3	nad4L	cob	nad5	nad2
------	------	------	------	------	------	------	-------	------	------	------	-------	-----	------	------

Plakortis simplex, *Plakortis halichondroides*, *Plakonarella cf. onkodes* DVL-2011, *Plakusa jani*, *Plakusa monophyla*, *Plakusa crypta*, *Plakusa* sp. DVL-2010 and *Corctisum candidabum* (Porifera):

DRNA	nad4	nad6	nad1	nad1	nad4L	nad2	nad5	cox1	cob	cox2	atp8	atp6	cox1	srRNA
------	------	------	------	------	-------	------	------	------	-----	------	------	------	------	-------

Branchiostoma floridae, *B. lanceolatum* and *Epigonostichus maldivensis* (Chordata: Cephalochordata), *Ejutatus burgeri* and *Myxine glutinosa* (Chordata: Craniata: Hyperotreti), *Homo sapiens*, *Mus musculus*, *Chelonia mydas*, *Typhlosceles natans*, *Xenopus laevis*, *Lattimeria chalumnae* and *L. madaonensis* (Chordata: Craniata: Vertebrata) etc. Chordates ancestral mitochondrial gene order:

nad1	nad2	cox1	cox2	atp8	atp6	cox3	nad3	nad4L	nad4	nad5	nad6	cob	srRNA	1rRNA
------	------	------	------	------	------	------	------	-------	------	------	------	-----	-------	-------

Strongylocentrotus purpuratus and *Paracentrotus lividus* (Echinodermata: Echinoidea), *Apostichopus japonicus* and *Cucumaria minima* (Echinodermata: Holothuroidea) etc. Echinoderms ancestral mitochondrial gene order:

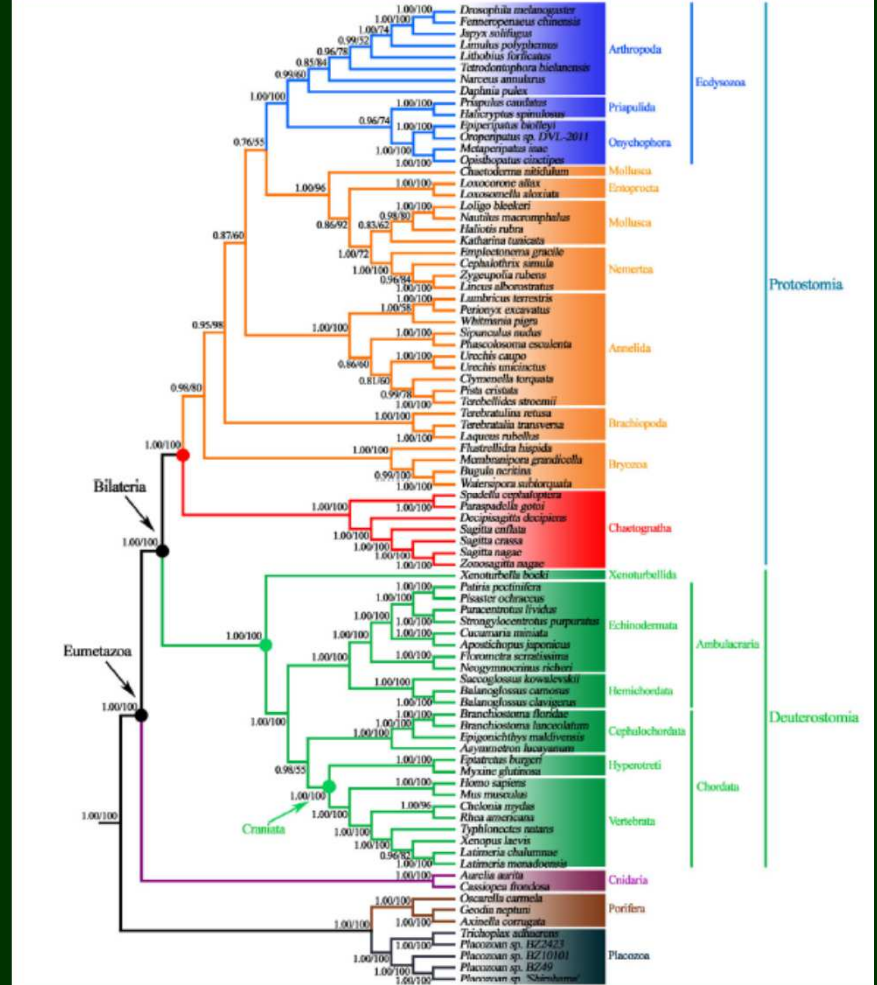
nad1	nad2	1rRNA	cox1	nad4L	cox2	atp8	atp6	cox3	nad3	nad4	nad6	cob	srRNA
------	------	-------	------	-------	------	------	------	------	------	------	------	-----	-------

Balanoglossus carnosus and *B. claviporus* (Hemichordata), Hemichordates ancestral mitochondrial gene order:

srRNA	DRNA	nad2	nad1	nad2	cox1	cox2	atp8	atp6	cox3	nad3	nad4L	nad4	nad5	cob	nad6
-------	------	------	------	------	------	------	------	------	------	------	-------	------	------	-----	------

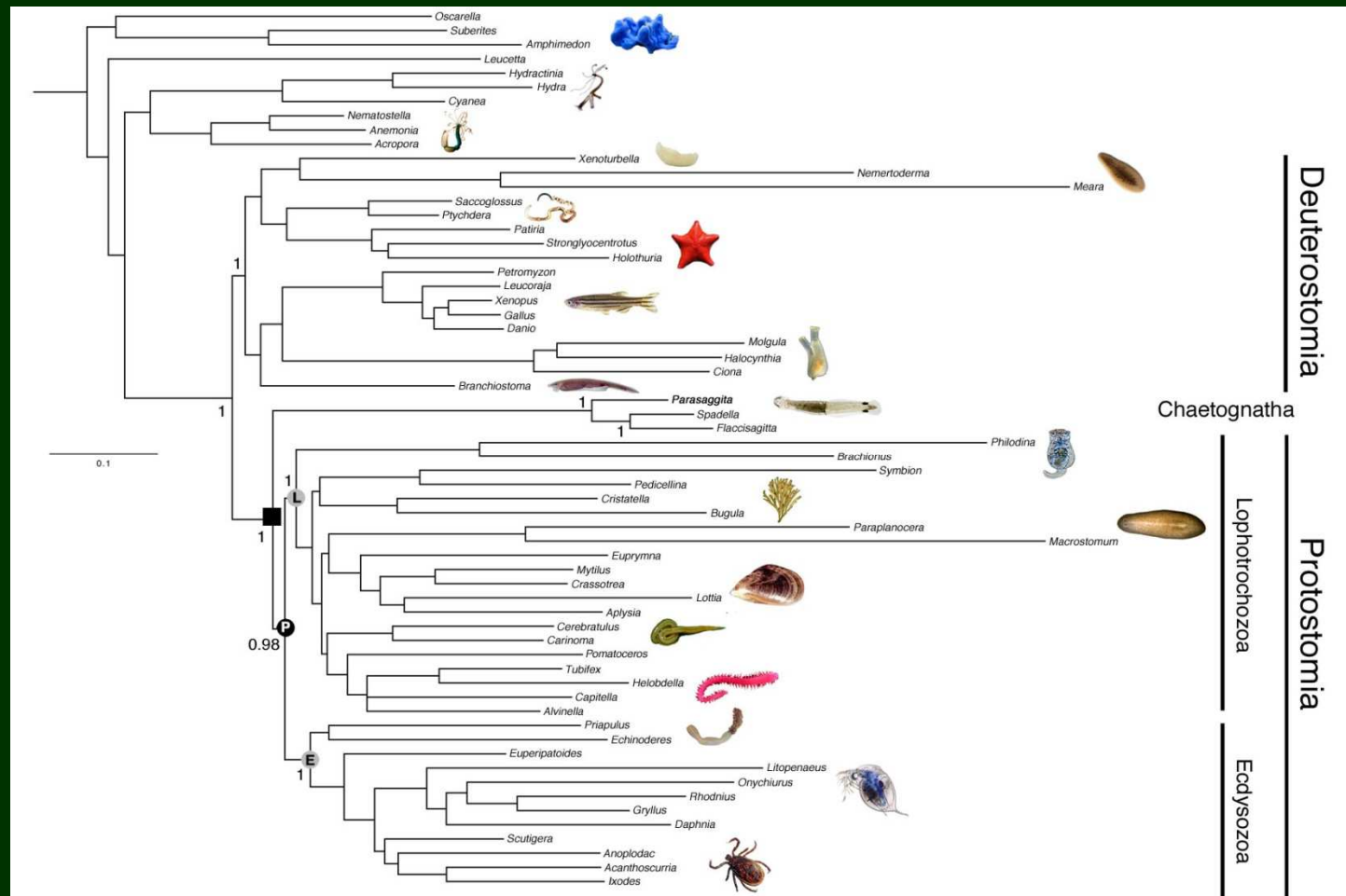
Xenoturbella bocki (Xenoturbellida):

nad1	nad2	cox1	cox2	atp8	atp6	cox3	nad3	nad4L	nad4	nad5	cob	srRNA	1rRNA	nad6
------	------	------	------	------	------	------	------	-------	------	------	-----	-------	-------	------



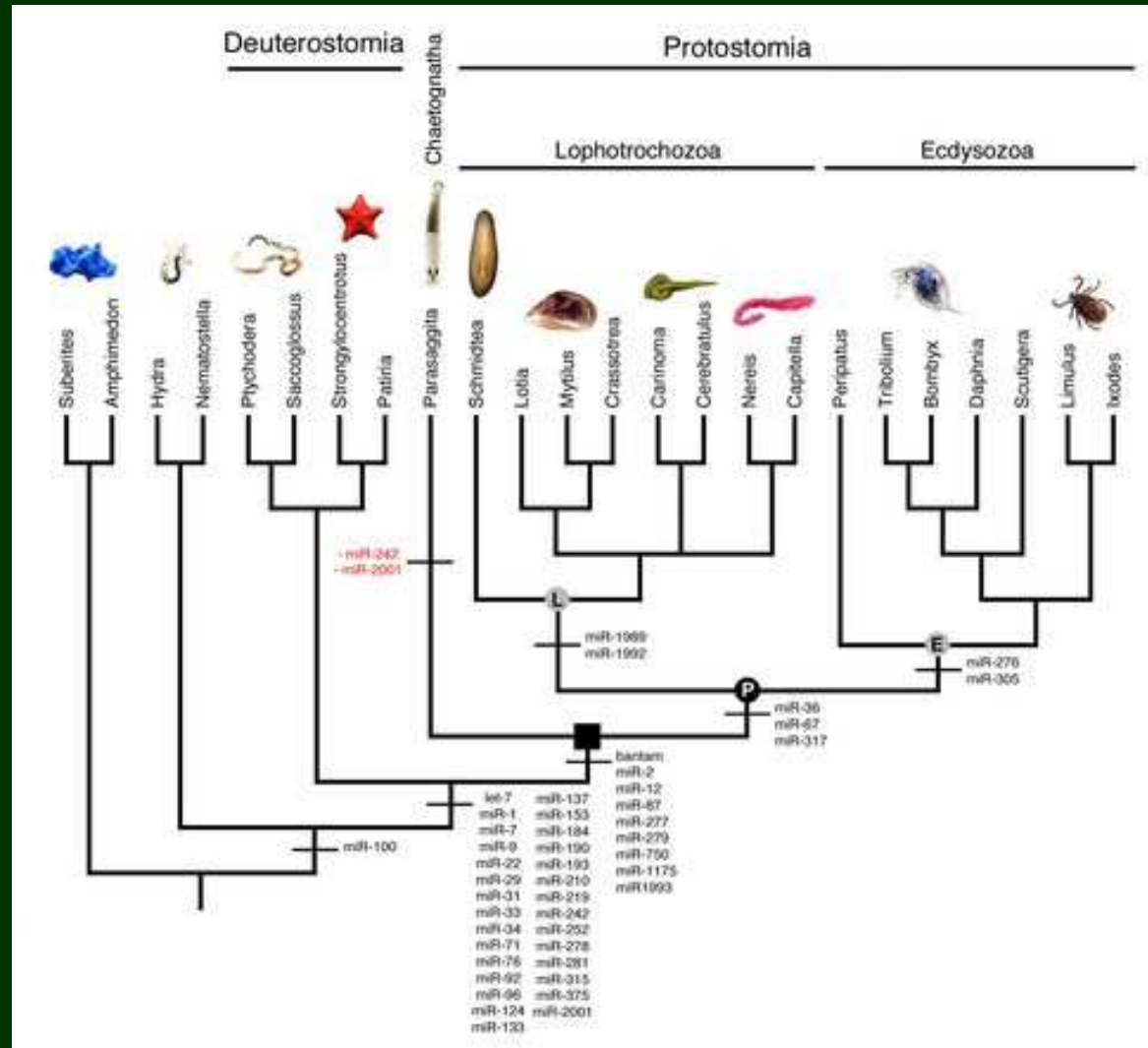
Fylogeneze ploutvenek

186 genu



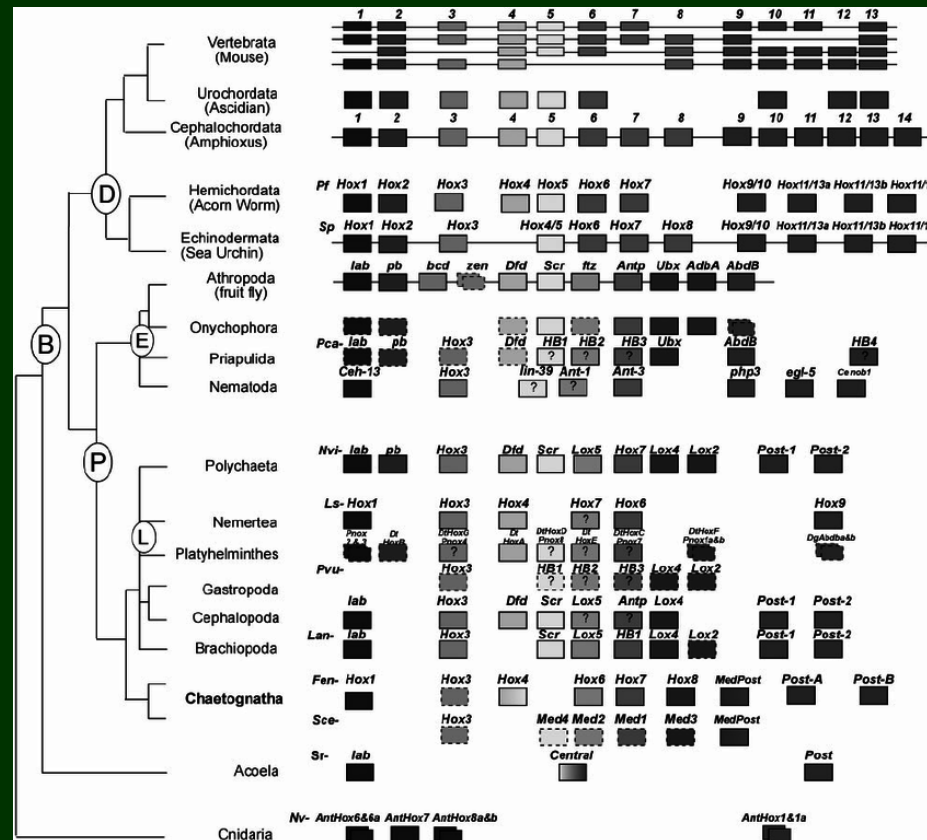
Fylogeneze ploutvenek

miRNA



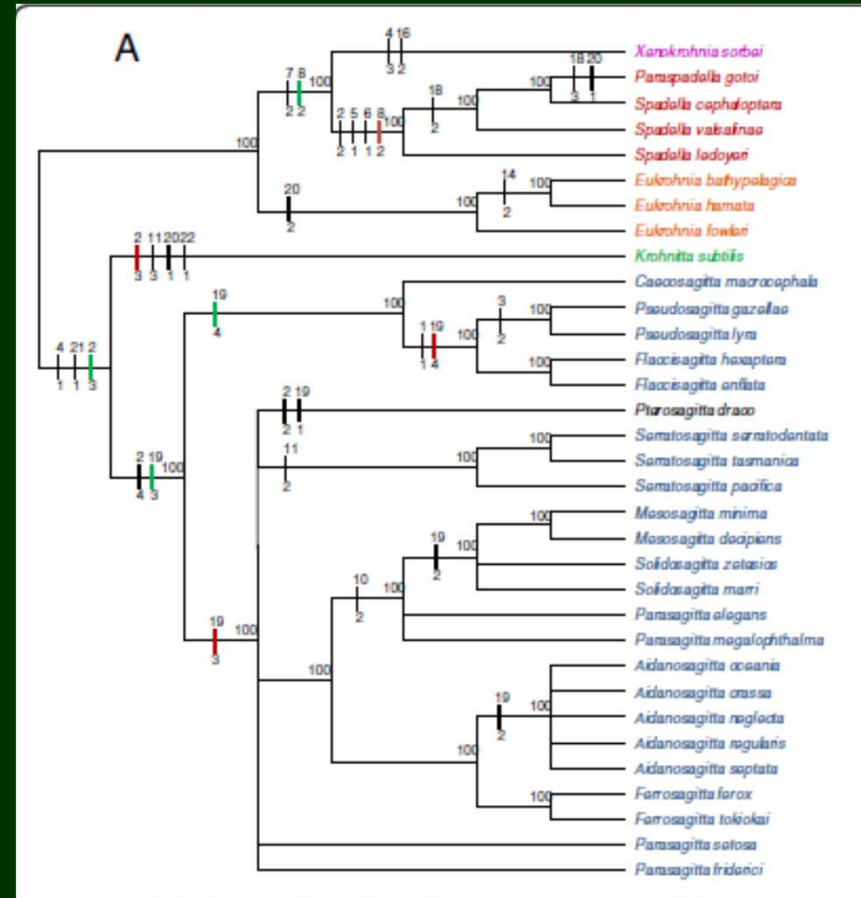
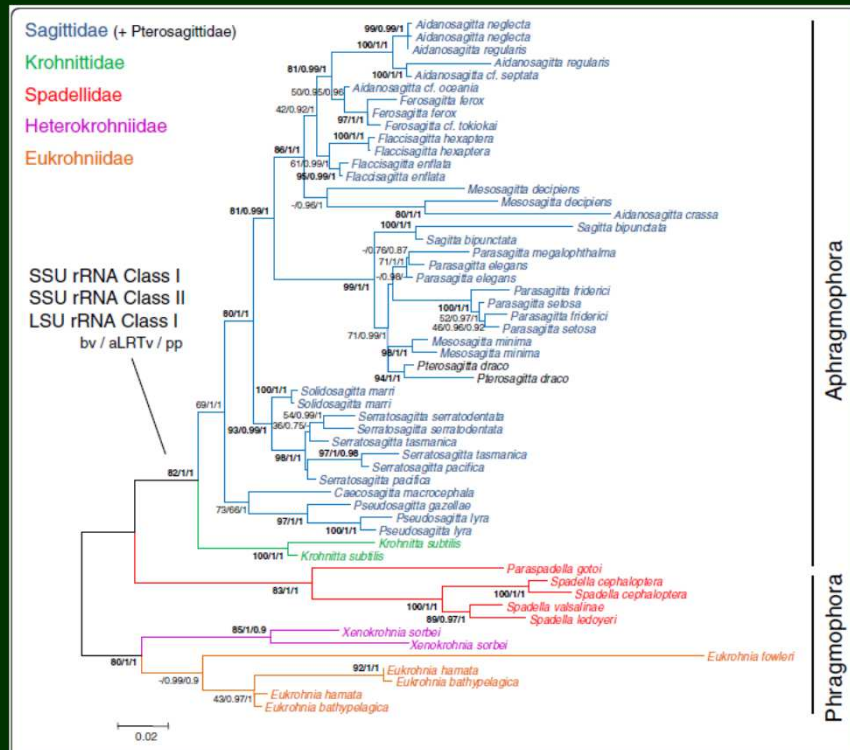
Chaetognatha Hox

- zvláštní (primitivní???) Hox geny – *MedPost* spojuje vlastnosti mediánních a posteriorních (ty nejsou příbuzné genům u spirálií a ekdysozoí)



Fylogeneze ploutvenek

18S + 28S + morfologie



Fylogeneze ploutvenek

- Phragmophora parafyletická (1. Eukrohniidae + Heterokrohniidae; 2. Spadellidae)
- Aphragmophora monophyletická
- převládající zjednodušení tělního plánu

