

Ctenophora Urochordata Nemertea
 Echinodermata Platyhelminthes
 Kamptozoa Nematomorpha
 Brachiopoda Porifera Phoronida
 Annelida Acoela Arthropoda
 Mollusca Bryozoa
 Onychophora Acanthocephala
 Nematoda Cnidaria Tardigrada
 Hemichordata

Ctenophora Urochordata Nemertea
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 Annelida Arthropoda
 Mollusca Bryozoa
 Cnidaria Acanthocephala
 Nematoda Hemichordata
 Rotifera

How are phyla related?

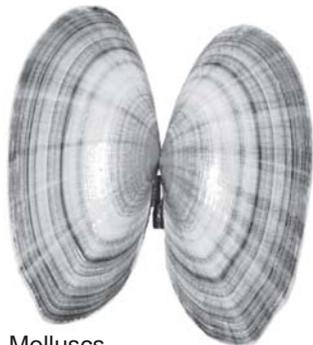
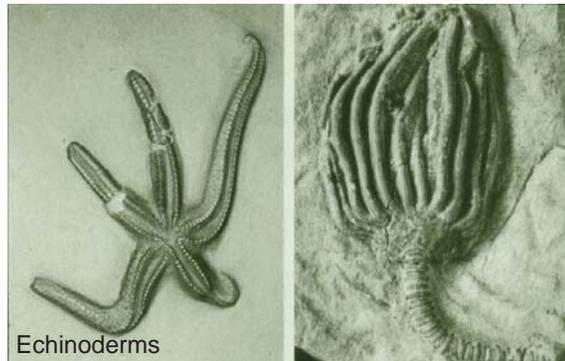
When did they appear?

How has diversity changed?

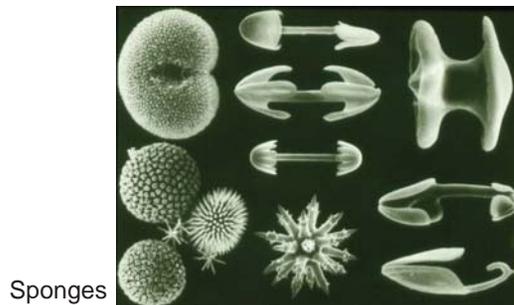
What mechanisms underlie diversification?

Variable preservation among taxa

Which phyla are more likely to leave informative records?

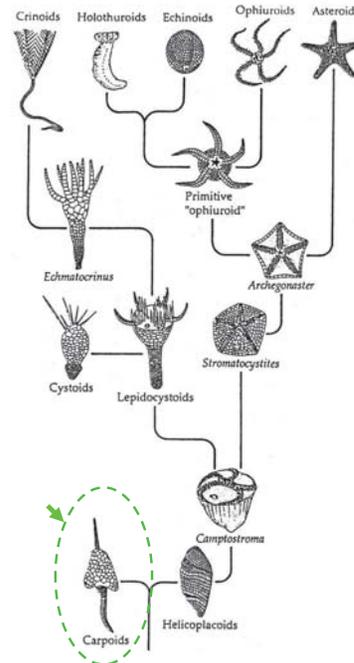


Molluscs

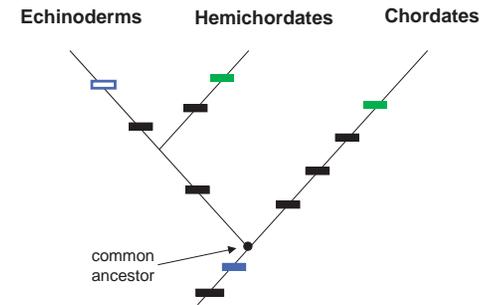


Sponges

1. Analysis of fossil taxa eg. Echinoderm classes



2a. Cladistics: phenotypes of extant taxa eg. Deuterostomes

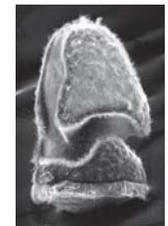


b. Cladistics: phenotypes of larval forms

(i) "dipleurula" type larva



holothuroid echinoderm auricularia

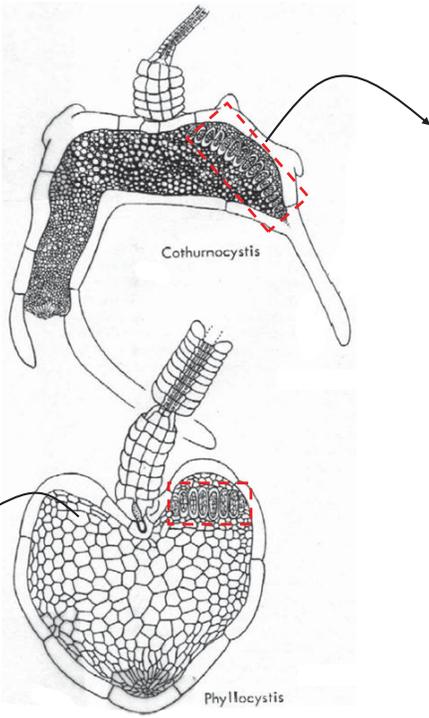


hemichordate tornaria

Carpoids



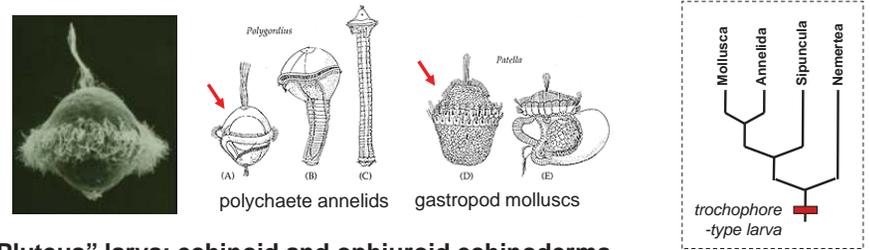
stereom



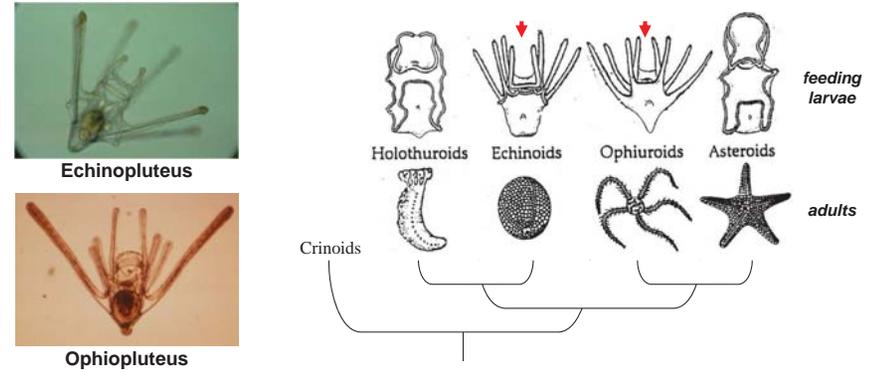
pharyngeal slits?

Are larval forms always reliable cladistic characters?

(ii) Trochophore-type larva: molluscs, annelids, sipunculans, nemerteans

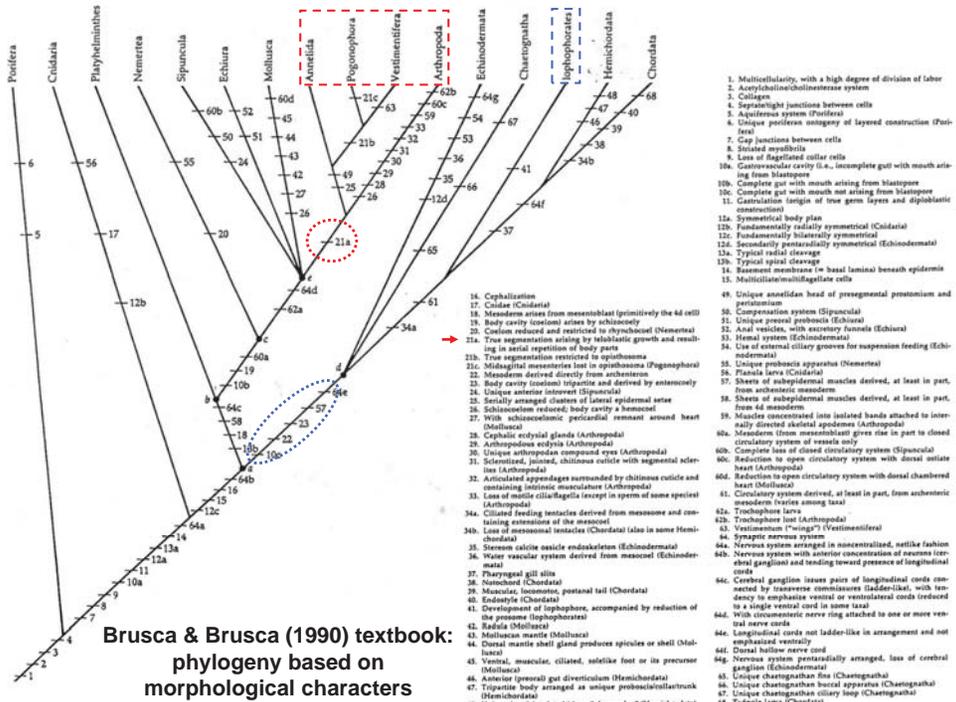
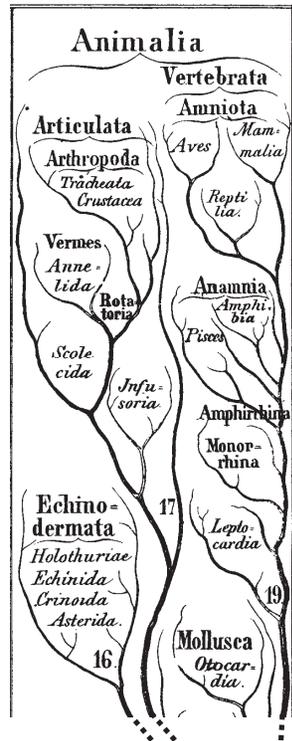
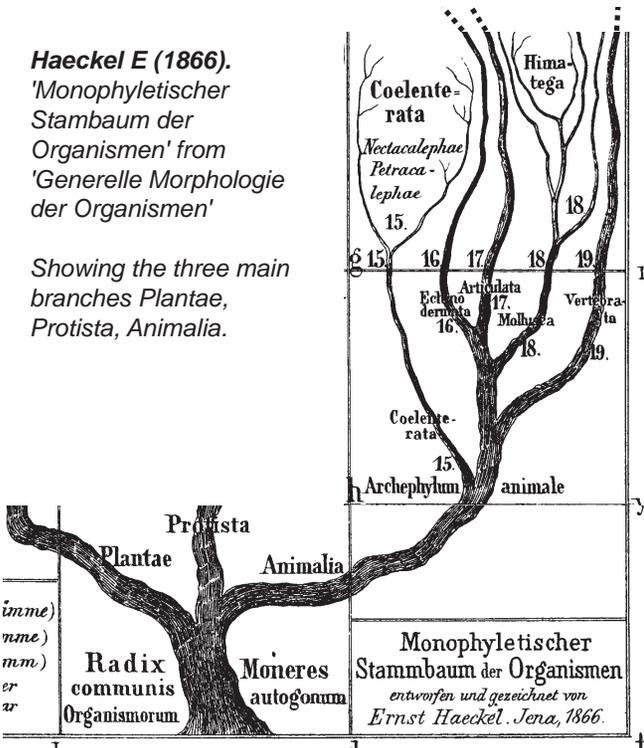


(iii) "Pluteus" larva: echinoid and ophiuroid echinoderms



Haeckel E (1866).
'Monophyletischer
Stambaum der
Organismen' from
'Generelle Morphologie
der Organismen'

Showing the three main
branches Plantae,
Protista, Animalia.

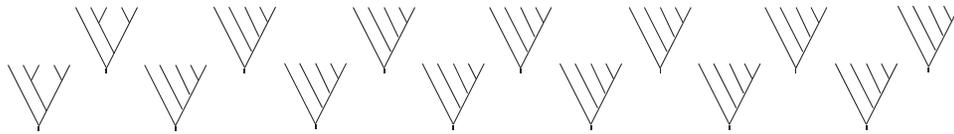
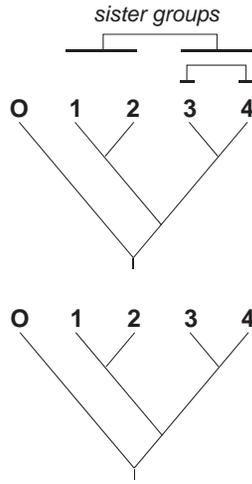


Brusca & Brusca (1990) textbook:
phylogeny based on
morphological characters

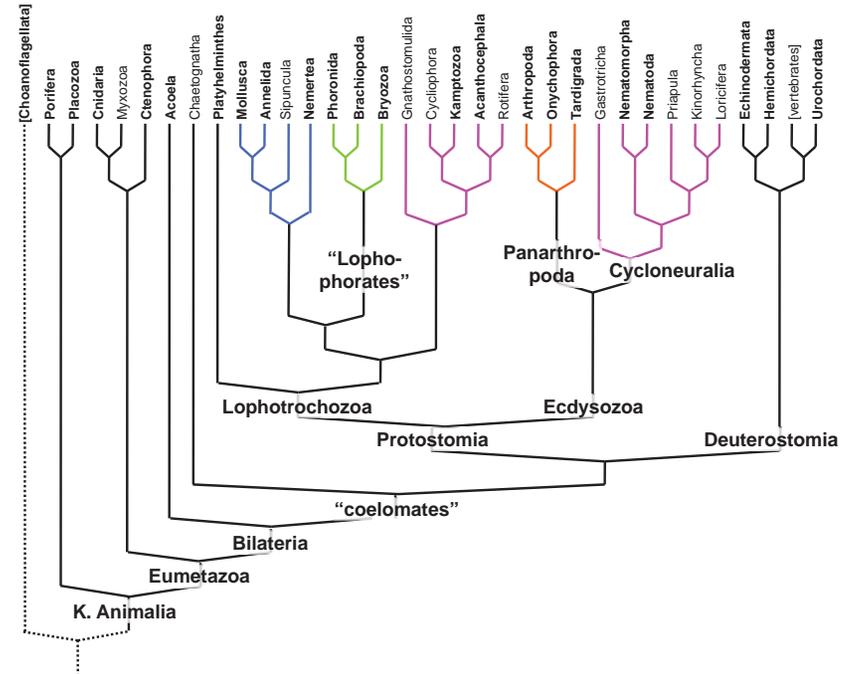
2b. Cladistics: molecular characters

DNA base	1	2	3	4	5	6	7	8	9	10	11	12
sp "O"	A	C	G	C	G	G	T	C	A	T	T	A
sp 1	.	G	T
sp 2	.	G	.	.	A	T
sp 3	.	.	.	T	.	C	.	.	.	A	.	T
sp 4	.	.	.	T	.	C	T

sp "O"	A	C	G	C	G	G	T	C	A	T	T	A
sp 1	.	G	T
sp 2	.	G	.	.	T	T
sp 3	.	.	.	T	.	C	.	.	.	A	.	T
sp 4	.	.	.	T	.	C	T

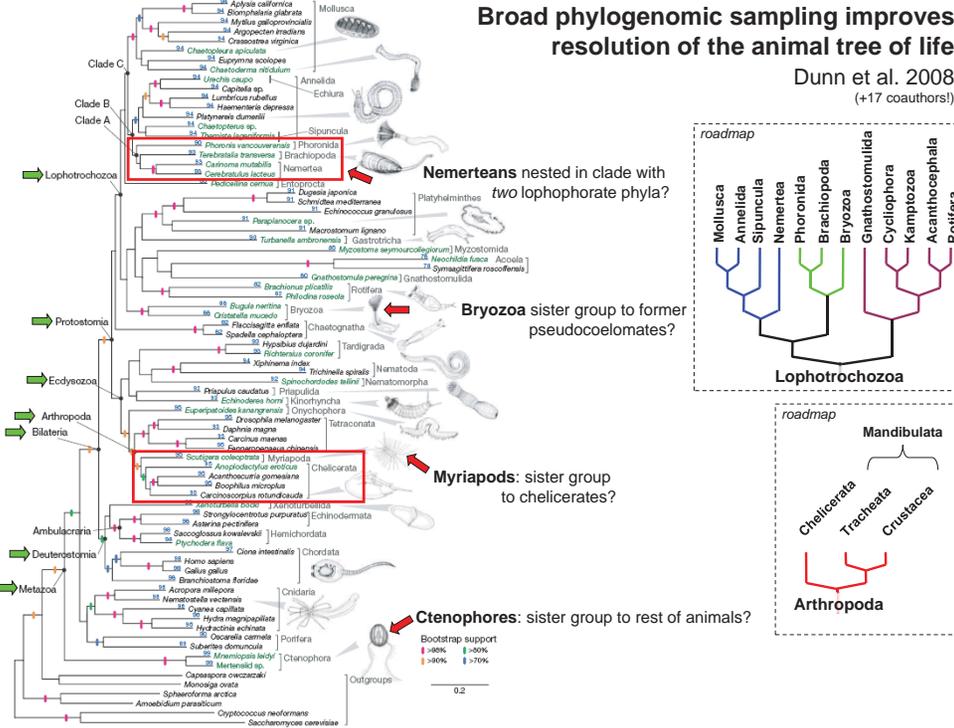


Our phylogenetic roadmap: a consensus view



Broad phylogenomic sampling improves resolution of the animal tree of life

Dunn et al. 2008 (+17 coauthors)



The Genome of the Ctenophore *Mnemiopsis leidyi* and Its Implications for Cell Type Evolution

Science (2013)

Joseph H. Ryan, Kevin Pang, Christian E. Schmitt, Ash-Dee Nguyen, R. Travis Moreland, David K. Slomovitz, Bharat J. Kish, Warren R. Francis, Paul Hradok, NISC Comparative Sequencing Program, Stephen A. Smith, Nicholas H. Putnam, Steven H. D. Bradlock, Casey W. Dunn, Ilya G. Webber, James C. Mullikin, Mark D. Martindale, Andreas D. Baxevanis

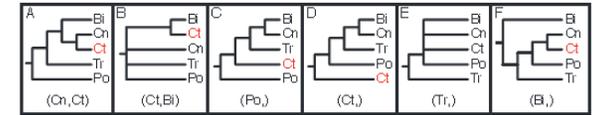
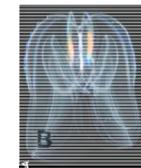
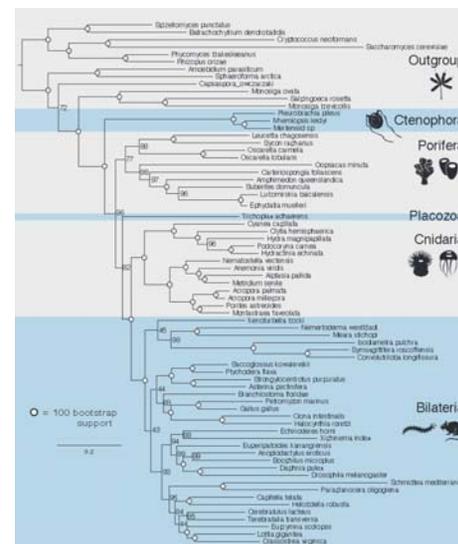
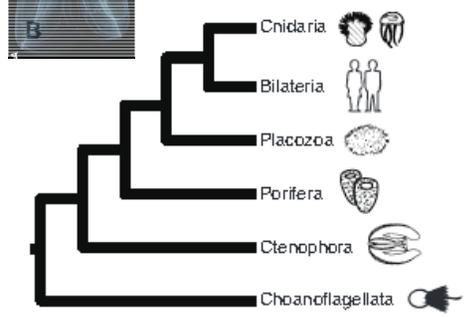


Fig. 2. Previously proposed relationships of the five deep clades of animals. The label at the bottom of each pane corresponds to the header of Table 1. (A) Coelenterata hypothesis, (E) Ctenophora as sister to Bilateria, (C) Porifera as sister group to the rest of Metazoa, (D) Ctenophora as sister group to the rest of Metazoa, (F) Diphlocladia hypothesis. We see no support in any of our analyses for the hypotheses in (A), (E), and (F) and very little support for (B) (see Table 1). (C), Ctenophora; Po, Porifera; Tr, Placozoa; Ch, Cnidaria; Bi, Bilateria.



The phylogenetic position of the ctenophore *Mnemiopsis leidyi* and its implications regarding the origin of mesodermal cell types. (A) Adult *M. leidyi*. (B) Summary of the relationships of the five main branches of animals and the outgroup Choanoflagellata.

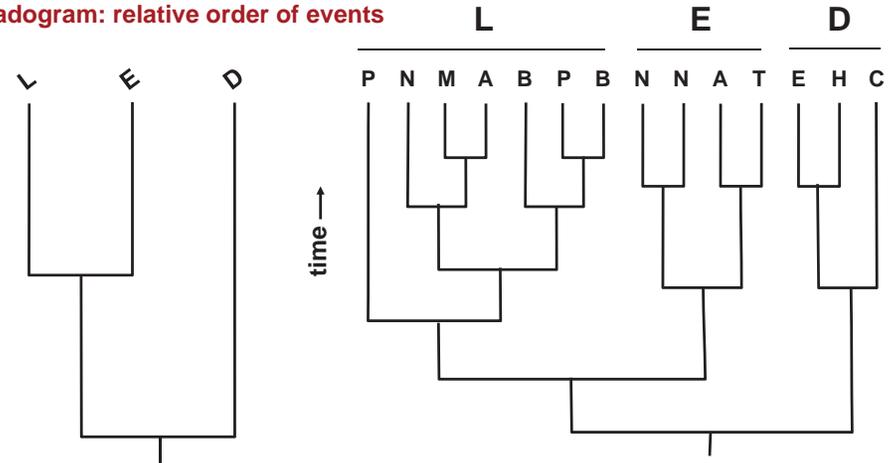


muscles? nerves? epithelia? axes?

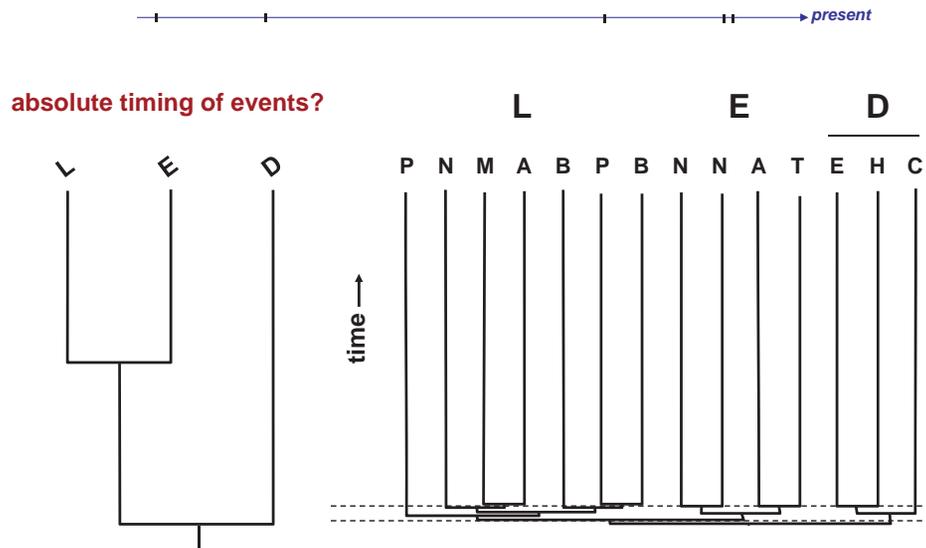
Pattern of evolutionary relationships

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How are phyla related?
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When did they appear?
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How has diversity changed?
 Annelida Acoela Anthropoda
What mechanisms underlie diversification?
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 Cnidaria Tardigrada
 Nematoda Hemichordata

cladogram: relative order of events



Timing of evolutionary divergences?



The major Pre-Cambrian and Cambrian Lagerstätten

Pre-Cambrian

Bitter Springs	1000–850 Ma	South Australia
Ediacara Hills	630-542 Ma	South Australia
Doushantuo Formation	600–555 Ma	Guizhou Province, China

Cambrian

Maotianshan Shales (Chengjiang)	525 Ma	Yunnan Province, China
Sirius Passet	518 Ma	Greenland
Emu Bay shale	517 Ma	South Australia
Kaili Formation	513–501 Ma	Guizhou province, south-west China
Wheeler Shale (House Range)	507 Ma	Western Utah, US
Burgess Shale	505 Ma	British Columbia, Canada
Kinnekulle Orsten and Alum Shale	500 Ma	Sweden
Öland Orste and Alum Shale	500 Ma	Sweden

Pre-Cambrian seas?



The Ediacaran (named after the hills in Australia, the site of a major pre-Cambrian fossil deposit) featured soft-bodied life - no bones, shells, teeth or other hard parts. Most of the life forms bear no resemblance to modern day animals, and may have become the eventually extinguished prey of more modern forms. The world's first ever burrowing animals evolved in the Ediacaran but left only trace fossils (burrows).

Enigmatic Ediacaran biota (630-542 mya)



Dickinsonia costata, an iconic Ediacaran organism, displays the characteristic quilted appearance of Ediacaran enigmata



Spriggina superficially resembled a segmented animal but the apparent segments are isomers (alternating on left and right sides).



Charniodiscus, probably a stationary filter feeder, was a frond anchored by a holdfast to a sandy sea bed.



Kimberella may have had a predatory or grazing lifestyle.



Archaeonassa-type trace fossils



Charnia, the first accepted complex Precambrian organism, once interpreted as a relative of the sea pens.

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Cambrian seas?

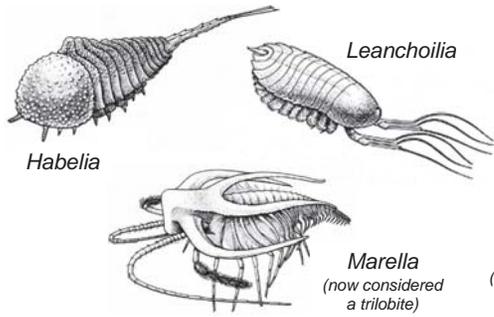


Some of the diversity of the Burgess Shale biota is depicted in the drawing above by Sam Gon III and John Whorral. Trilobites such as *Olenoides serratus* (1) were a minority among a diversity of arthropods such as *Sidneyia* (9), *Waptia* (17), *Helmetia* (13), *Sanctacaris* (18), *Tegopelte* (15), *Naraoia* (16), *Leanochoilia* (10), *Canadaspis* (12), *Odaraia* (19), *Marrella* (11), and *Burgessia* (14), as well as oddities such as *Opabinia* (24), *Wiwaxia* (26), *Hallucigenia* (20), and the giant predator, *Anomalocaris* (28).

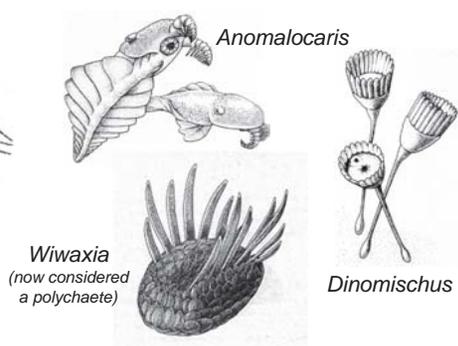
Problematic taxa of the Burgess Shale (525-505 mya)

Wonderful Life, Gould (1990)

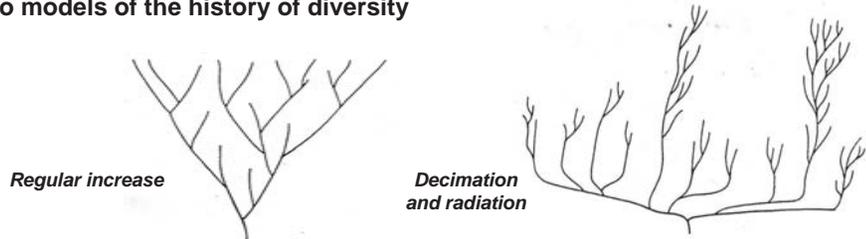
Unique arthropod subphyla?



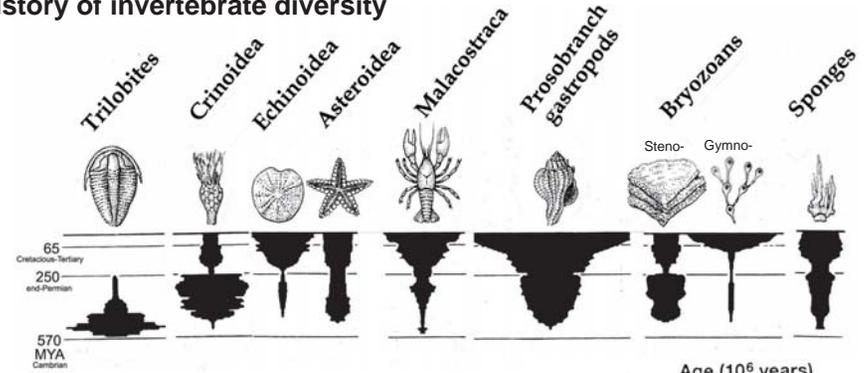
Unique phyla?



Two models of the history of diversity



History of invertebrate diversity



Mass extinctions of marine families

