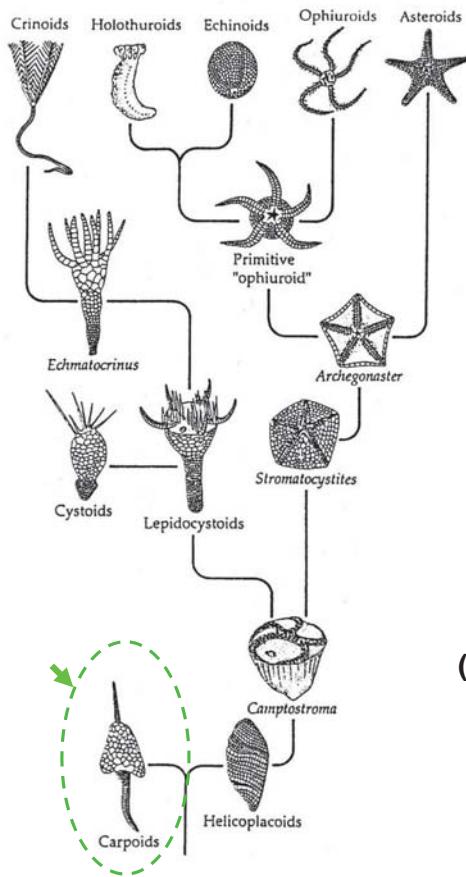
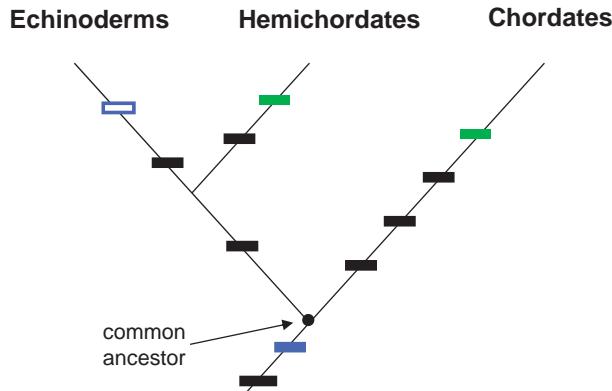


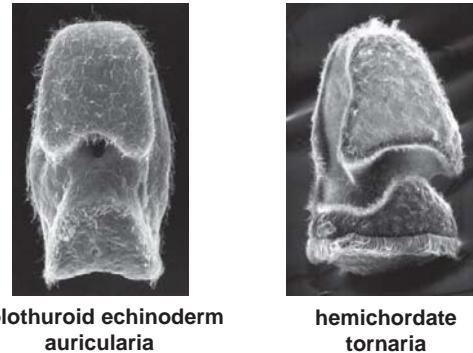
1. Analysis of fossil taxa eg. Echinoderm classes



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

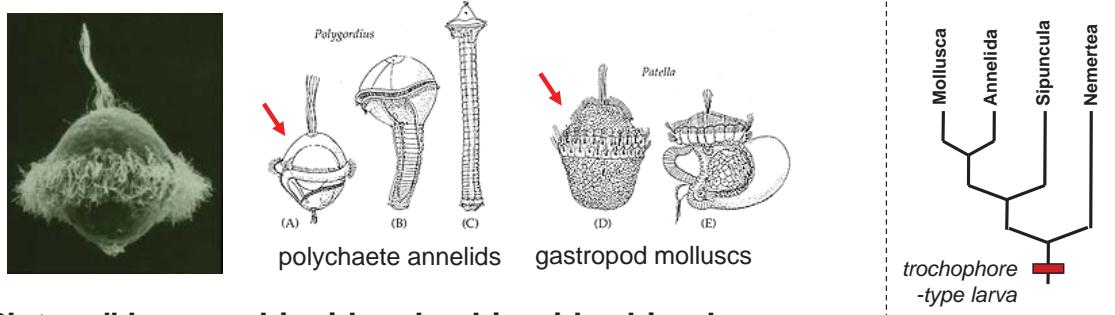


b. Cladistics: phenotypes of larval forms

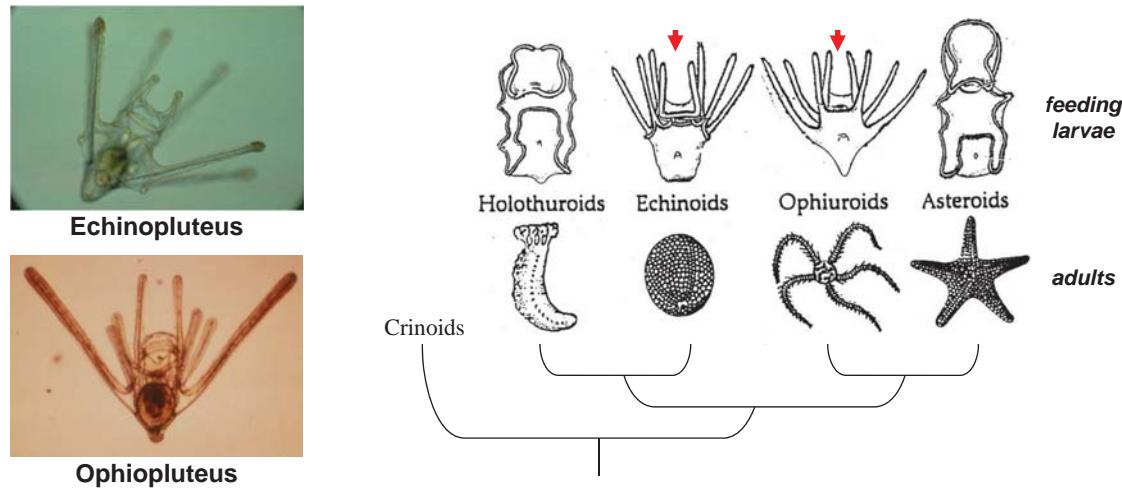


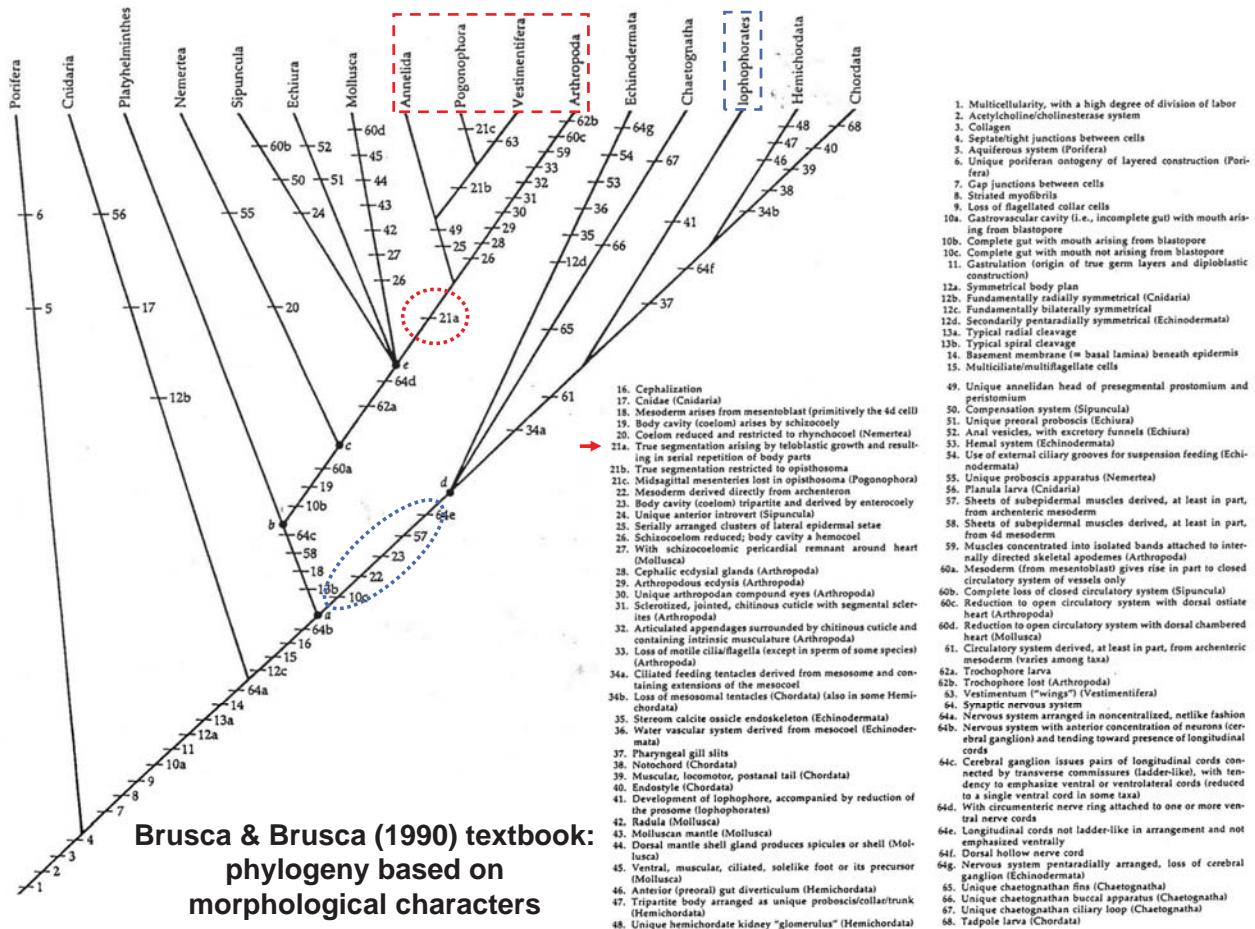
Are larval forms always reliable cladistic characters?

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



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

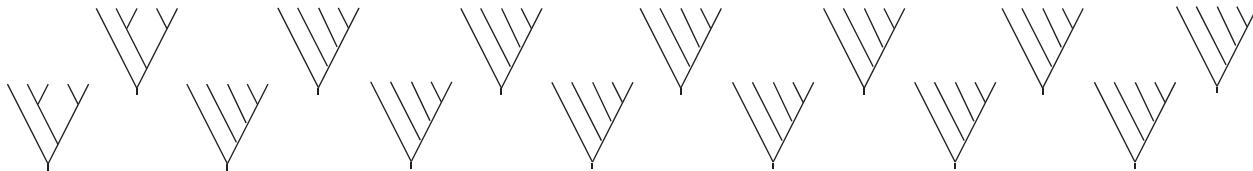
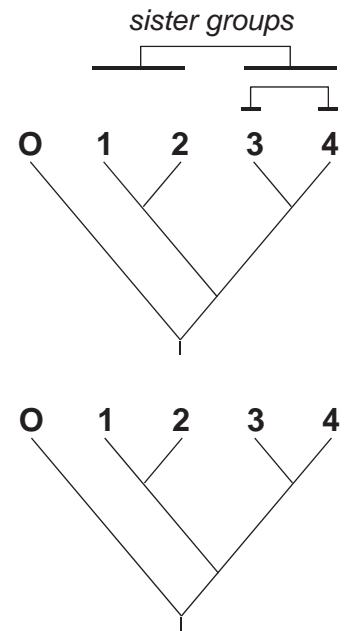




2b. Cladistics: molecular characters

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

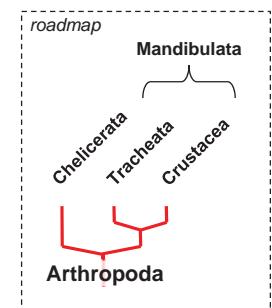
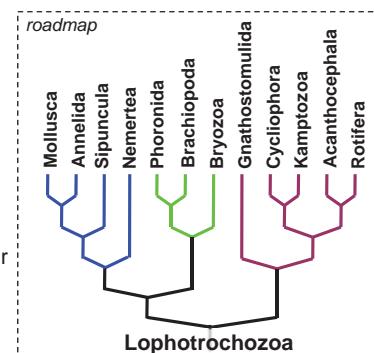
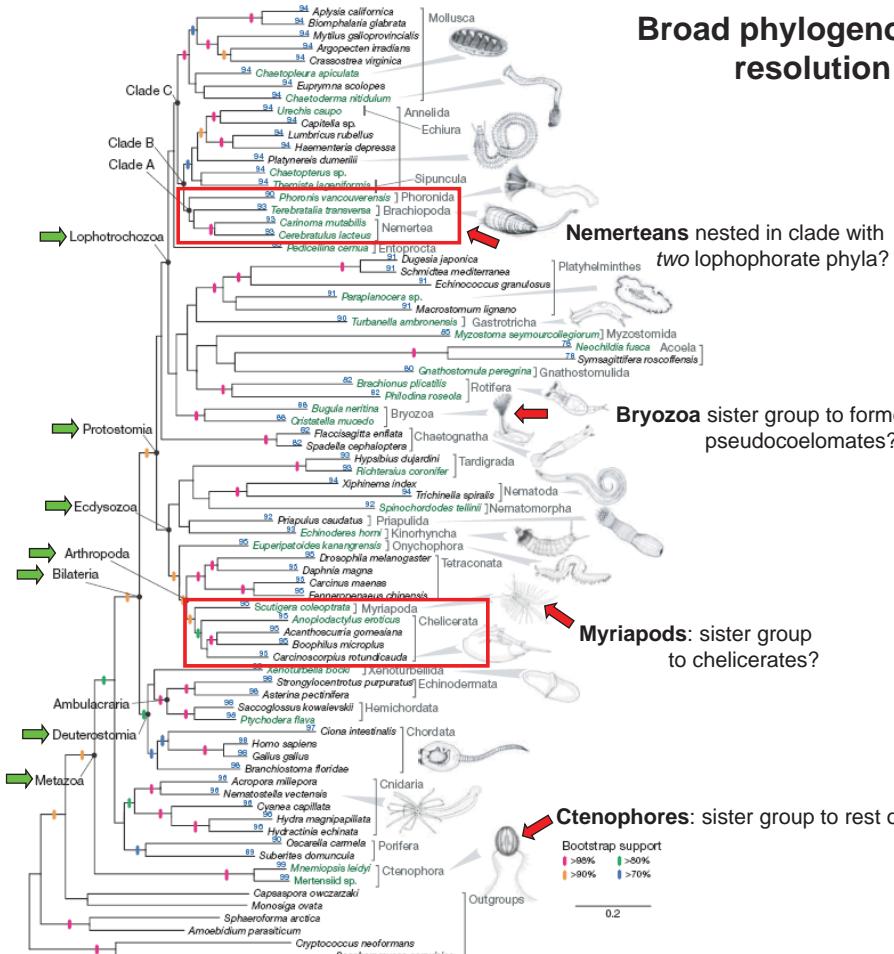
DNA base	1	2	3	4	5	6	7	8	9	10	11	12
sp "O"	A	C	G	C	G	T	C	A	T	T	A	
sp 1	.	G	T		
sp 2	.	G	.	.	T	T		
sp 3	T	.	C	.	A	.	T	
sp 4	T	.	C	.	.	.	T	



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

Joseph F. Ryan, Kevin Pang, Christine E. Schnitzler, Anh-Dao Nguyen, R. Travis Moreland, David K. Simlons, Bernard J. Koch, Warren R. Francis, Paul Huyak, NISC Comparative Sequencing Program, Stephen A. Smith, Nicholas H. Putnam, Steven H. D. Haddock, Casey W. Dunn, Tyra G. Wolfsberg, James C. Mullikin, Mark Q. Martindale, Andreas D. Baxevanis*

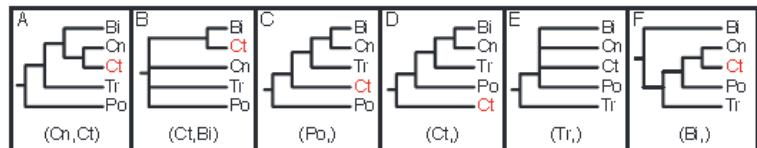
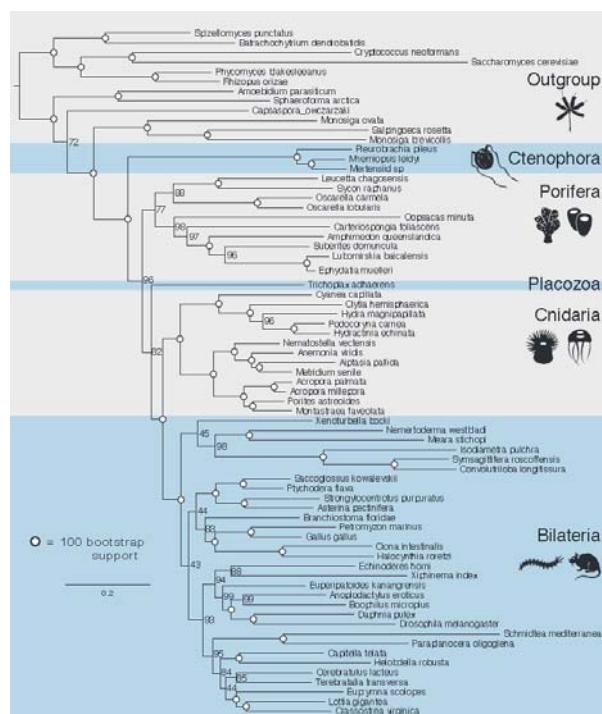
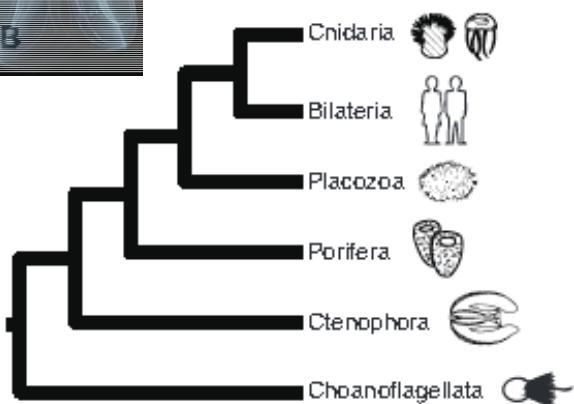


Fig. 2. Previously proposed relationships of the five deep clades of animals. The label at the bottom of each pane corresponds to the header of Table 1. (A) Coelenterata hypothesis. (B) Ctenophora as sister to Bilateria. (C) Porifera as sister group to the rest of Metazoa. (D) Ctenophora as sister group to the rest of Metazoa. (E) Placozoa as sister group to the rest of Metazoa. (F) Diploblastica 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). Ct, Ctenophora; Po, Porifera; Tr, Placozoa; Cn, Cnidaria; Bi, Bilateria.



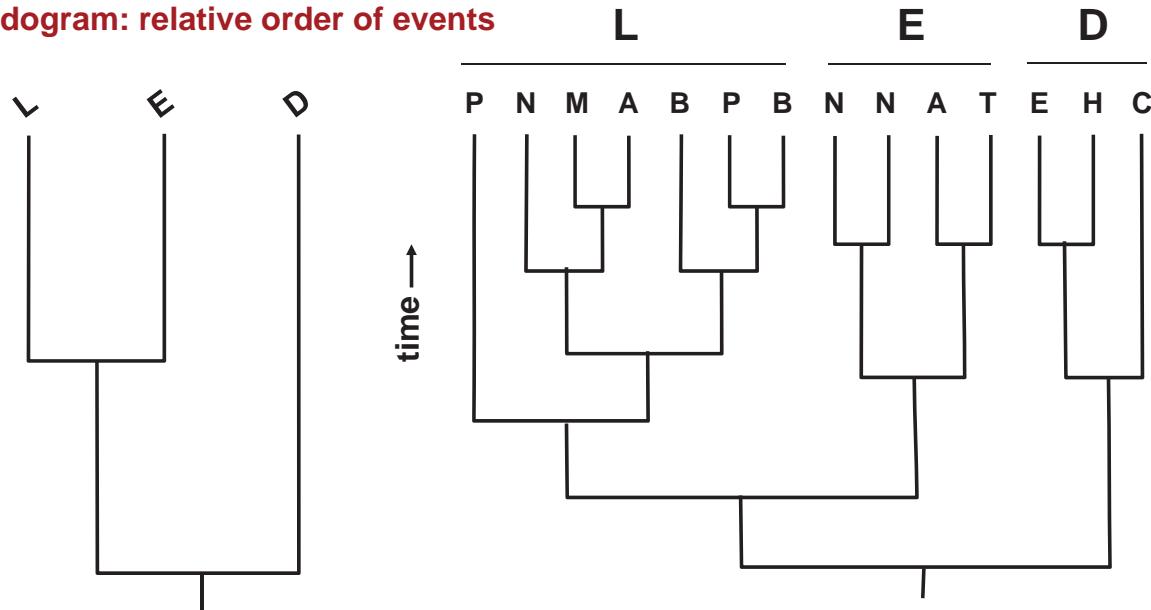
The phylogenetic position of the ctenophore *Mnemiopsis leidyi* and its implications regarding the origin of mesodermal cell types. (A) Adult *M. leidyi*. (B) Summary of the relationships of the five main branches of animals and the outgroup Choanoflagellata.



muscles? nerves? epithelia? axes?

Pattern of evolutionary relationships

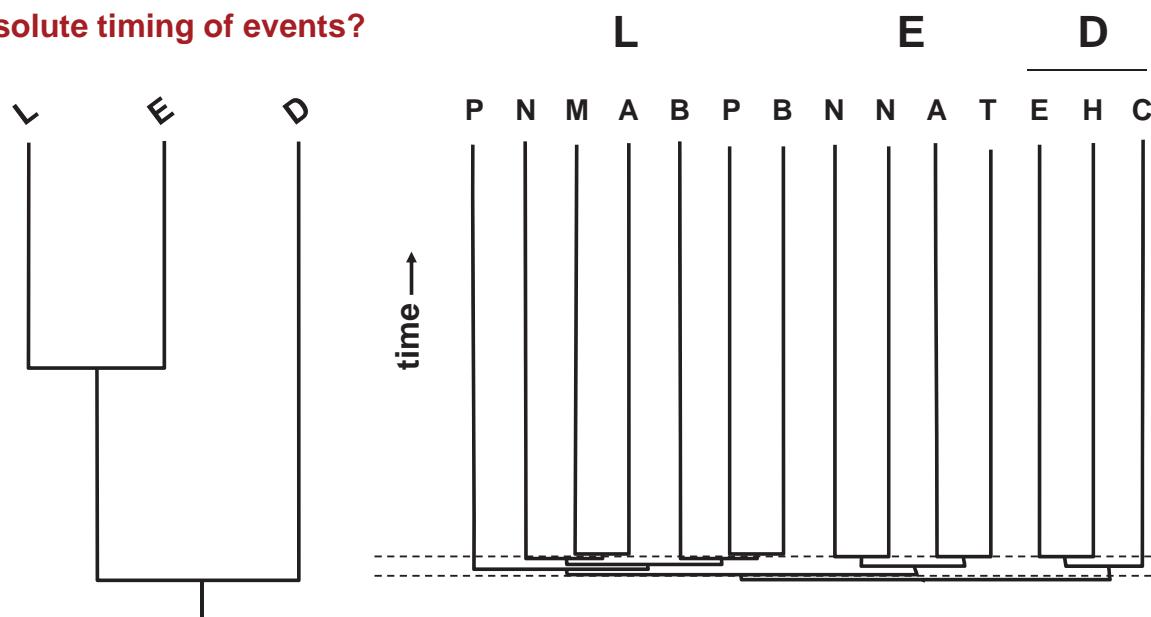
cladogram: relative order of events



Timing of evolutionary divergences?

present

absolute timing of events?



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

History of invertebrate diversity

