

# Multiple phyla, one time resolution?

## Time averaging in benthic foraminifera, mollusk, echinoid, crustacean and otolith fossil assemblages

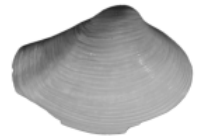
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universität  
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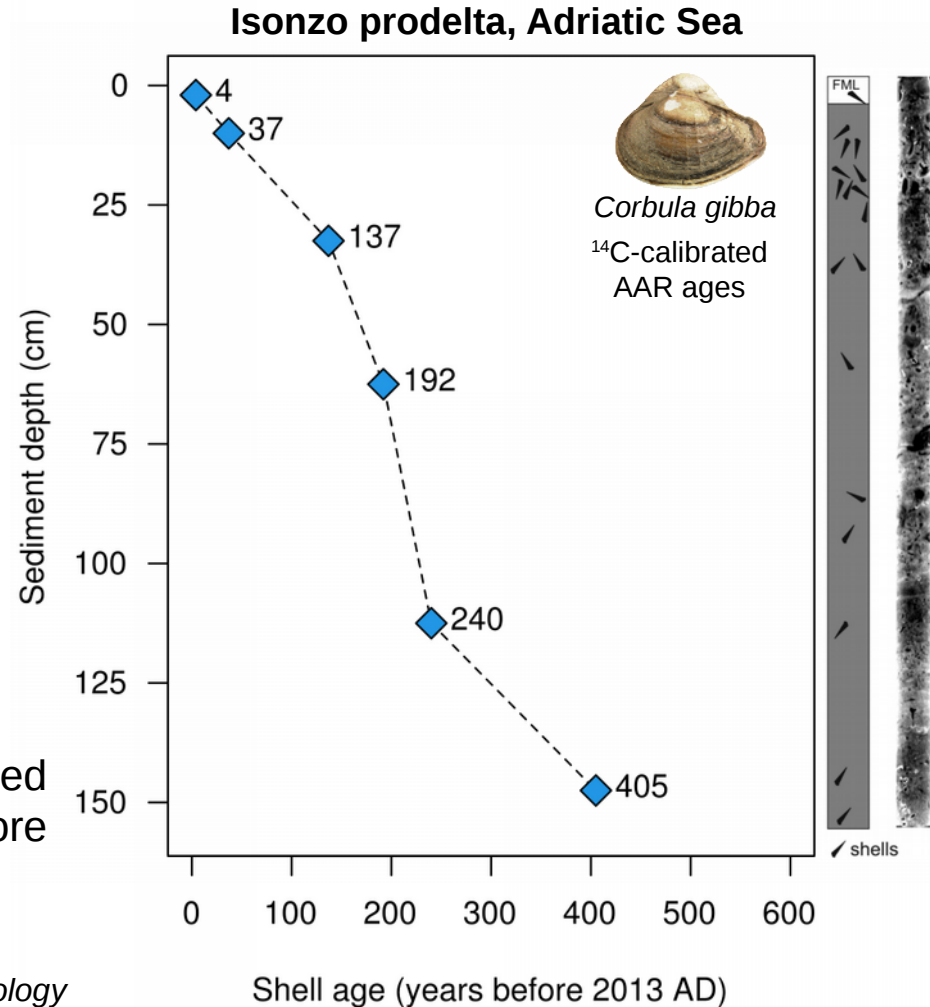


# Time averaging: the fundamental property of fossil assemblages

Fossil assemblages are frequently affected by extensive **time averaging**  
→ mixing of remains of organisms  
**separated by decades, centuries or millennia**  
in a single sedimentary layer

Time averaging limits the temporal resolution of **geochemical and geochronological records** based on biomineralized skeletal remains

6 shells dated  
in the entire core

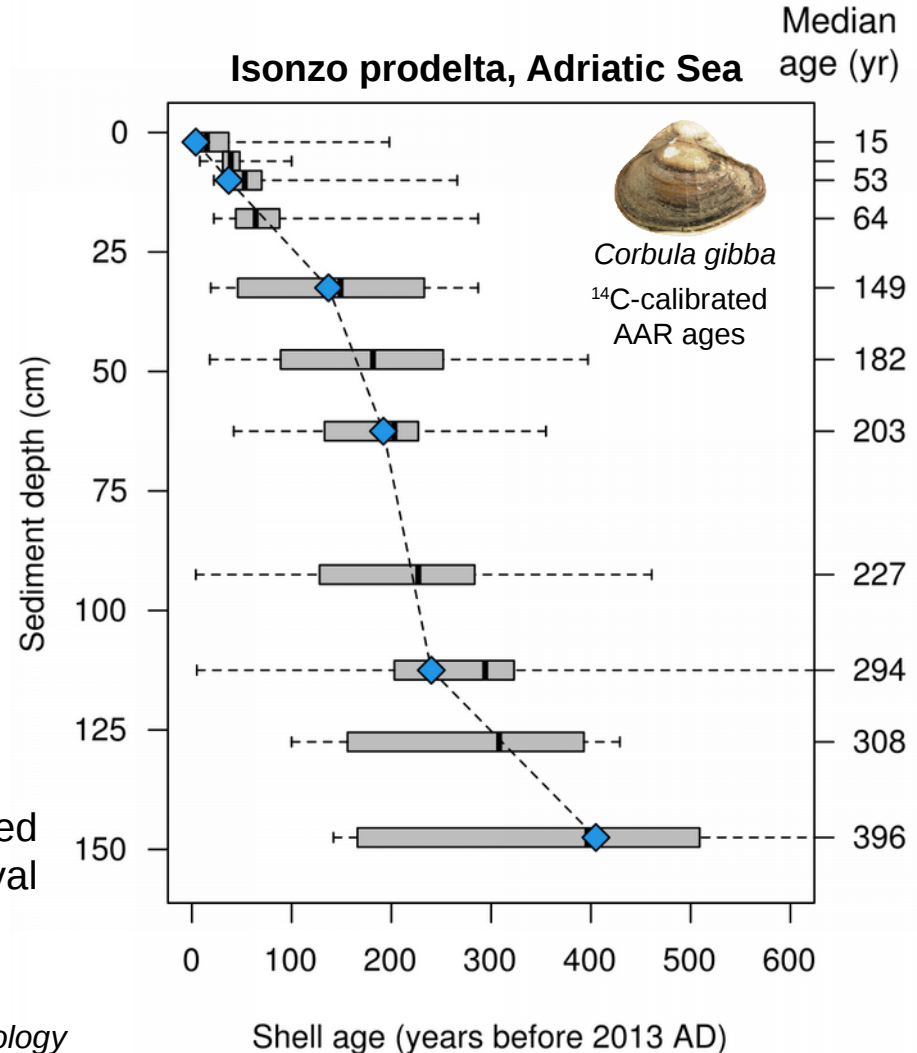


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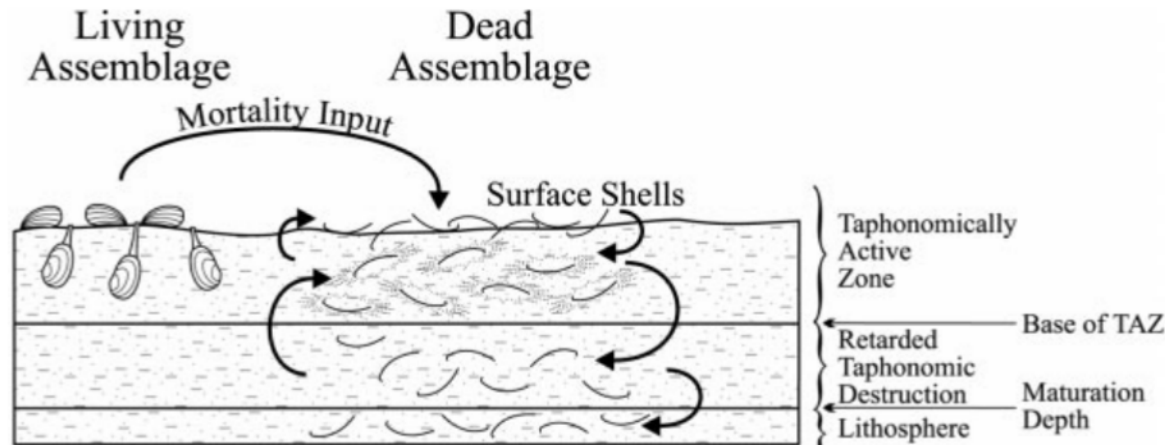
~30 shells dated  
per 5-cm core interval



# Time averaging: the fundamental property of fossil assemblages

What is controlling the variation in time averaging  
across taxa and environments?

- skeletal production rates
- skeletal disintegration rates
- sedimentation rates
- rate and depth of sediment mixing



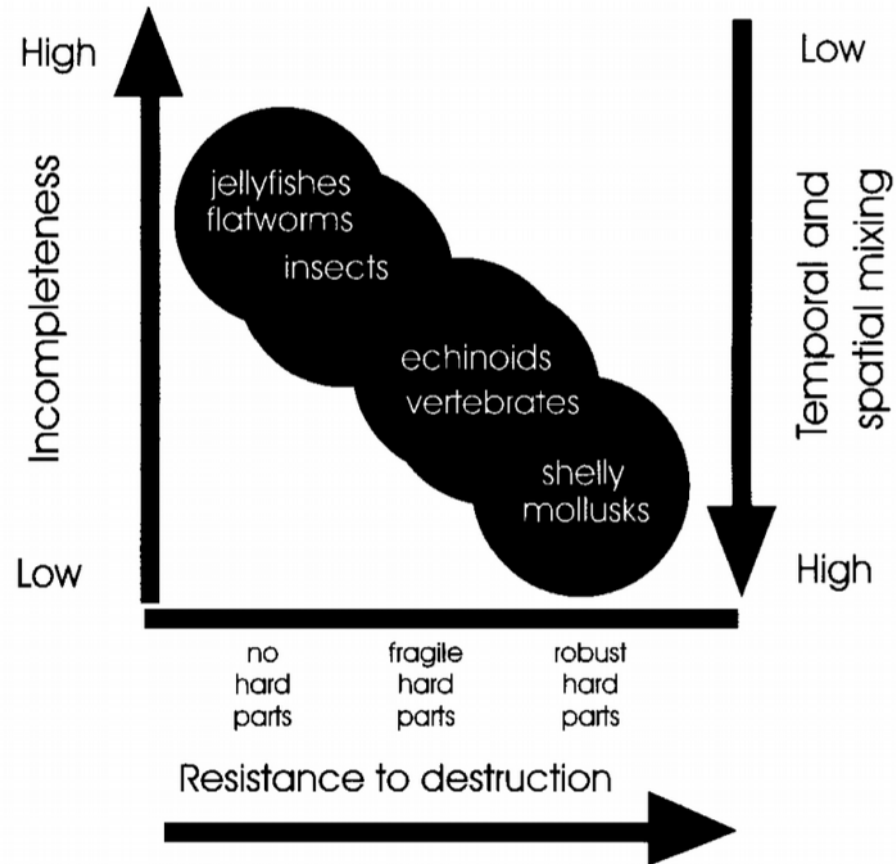
# Taxonomic variation in time averaging

Variation in skeletal durability across higher taxa may be the primary factor controlling the temporal resolution of fossil assemblages

More robust and larger remains should survive prolonged mixing near the sediment surface → higher time averaging

## How general in this pattern?

- many higher taxa lack direct estimates of time averaging
- processes that lead to burial and early diagenetic stabilization can allow fragile remains to escape rapid disintegration near the sediment-water interface

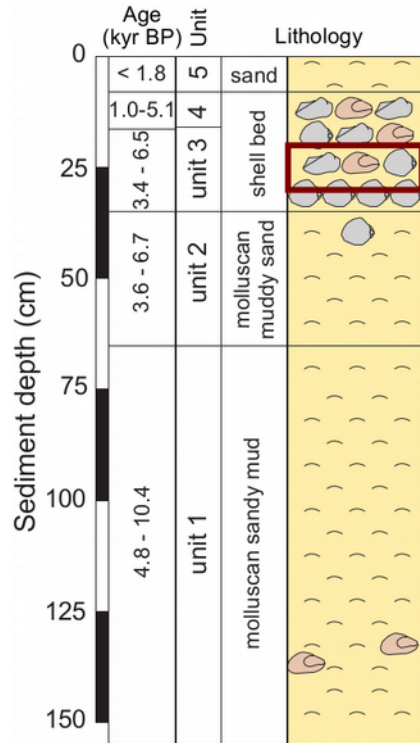


# Case study: time averaging in five phyla of carbonate producers

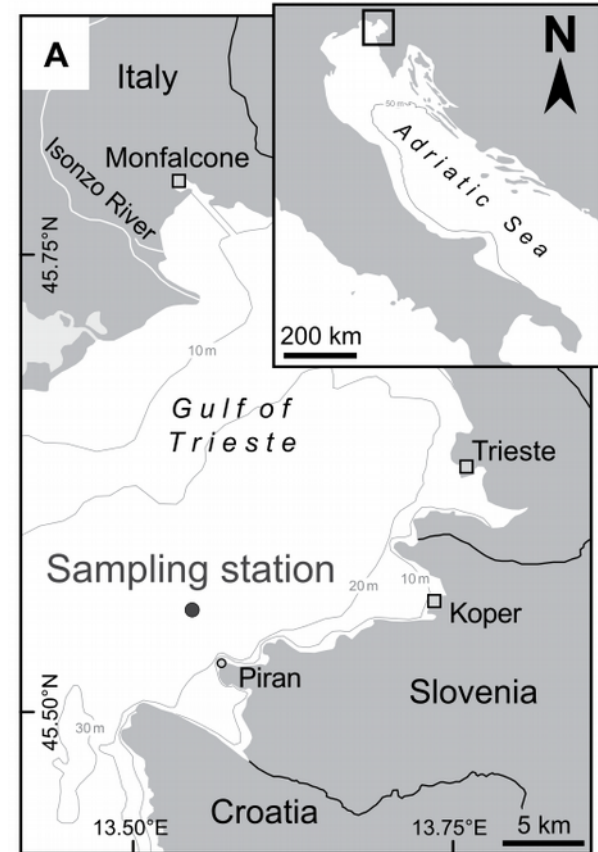
Sediment core from the NE Adriatic shelf (low sedimentation rate:  $\sim 0.15$  mm/yr)

Remains of mollusks, foraminifera, echinoderms, crustacean, and fish otoliths sampled from 10-cm subsurface shell bed interval

Age-dated with  $^{14}\text{C}$ -calibrated amino acid racemization (bivalves) and carbonate-target AMS  $^{14}\text{C}$  (all other taxa)



Station 4 km off Piran, Slovenia  
at 22.7 m water depth



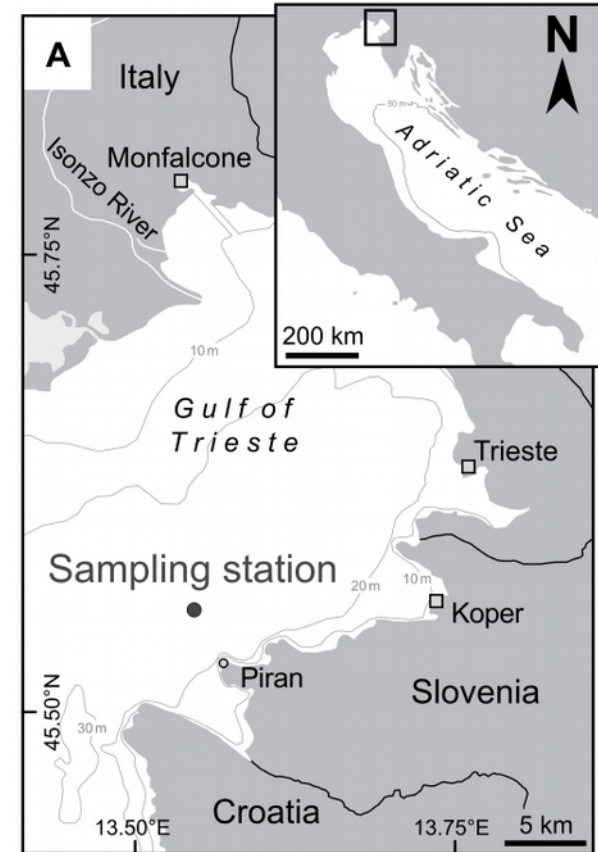
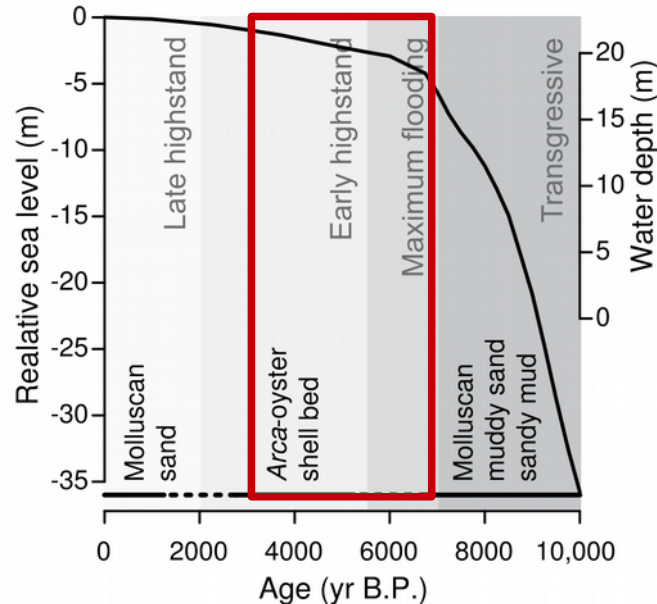
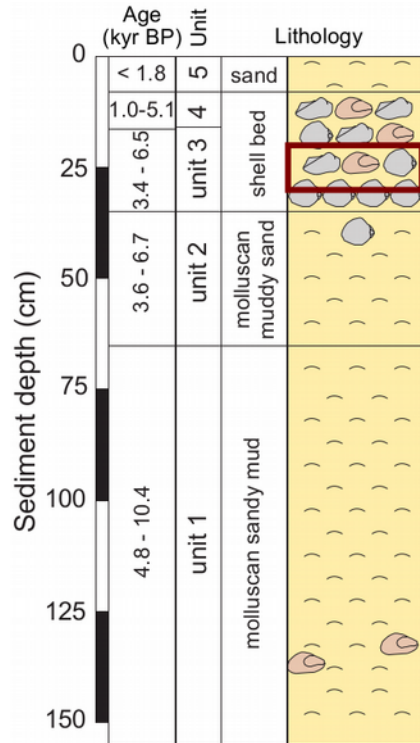


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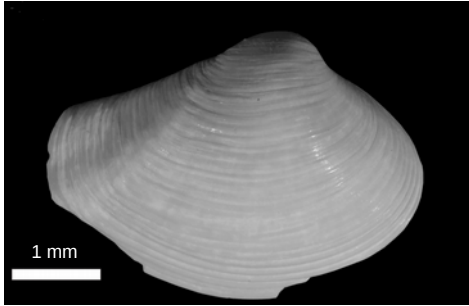


# Target taxa: different size, skeletal architecture and mineralogy

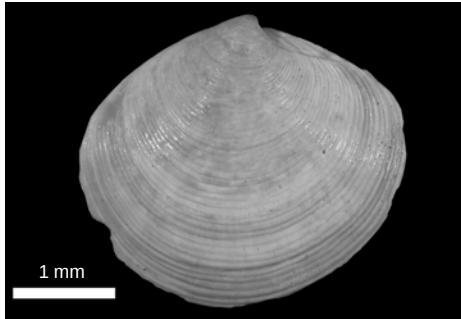
## Bivalves

aragonitic shells

*Corbula gibba*  
(Corbulidea)



*Gouldia minima*  
(Veneridae)



## Foraminifera

high- and low-Mg calcitic tests

*Adelosina intricata*  
(Miliolida, porcelaneous)



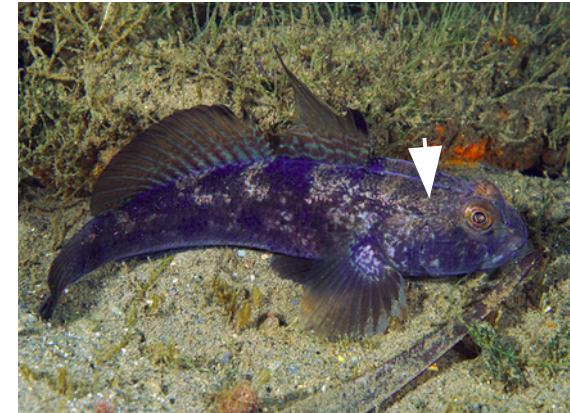
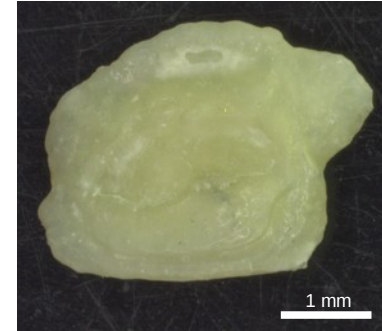
*Elphidium crispum*  
(Rotaliida, hyaline)



## Teleost fish

aragonitic otoliths

*Gobius cobitis*  
(Gobiidae)



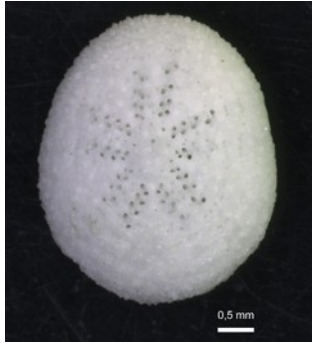


# Target taxa: different size, skeletal architecture and mineralogy

## Echinoids

high-Mg calcitic tests and madreporite plates

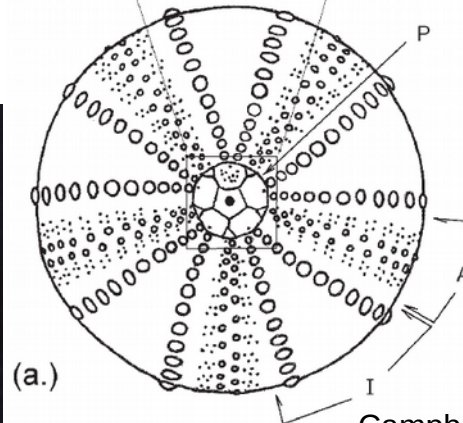
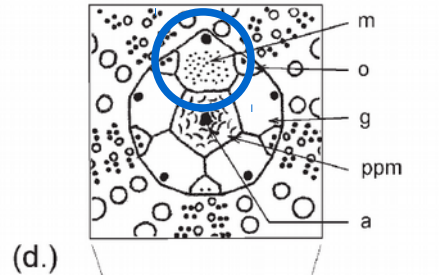
*Echinocyamus pusillus*  
(irregular clypeasteroid)



*Psammechinus* and *Paracentrotus*  
(regular camarodonts)



Madeporite  
(unique plate in echinoid test)

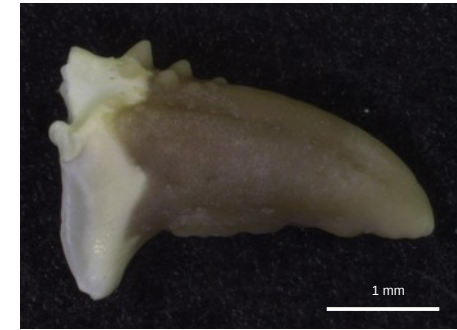


Campbell 2008

## Brachyuran crustacean

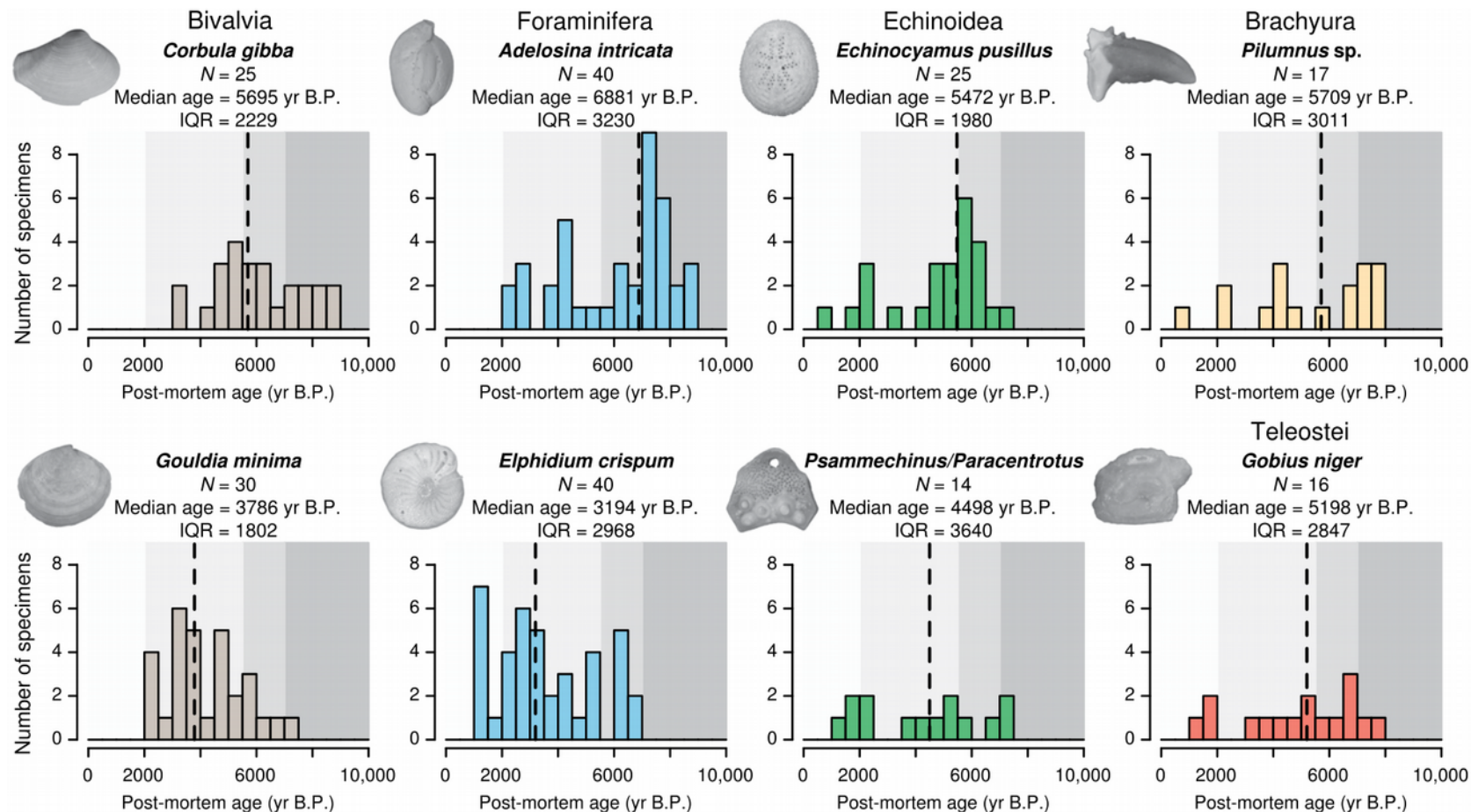
low-Mg calcitic chelae

*Pilumnus* sp.  
(Pilumnidae)



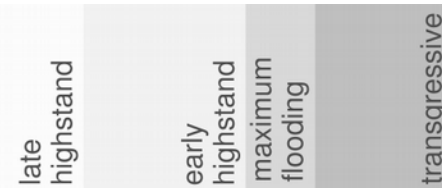
# Results: multiple phyla, one time resolution

Similar extent of time averaging in all taxa: ~1800 to ~3600 yrs (interquartile age ranges)

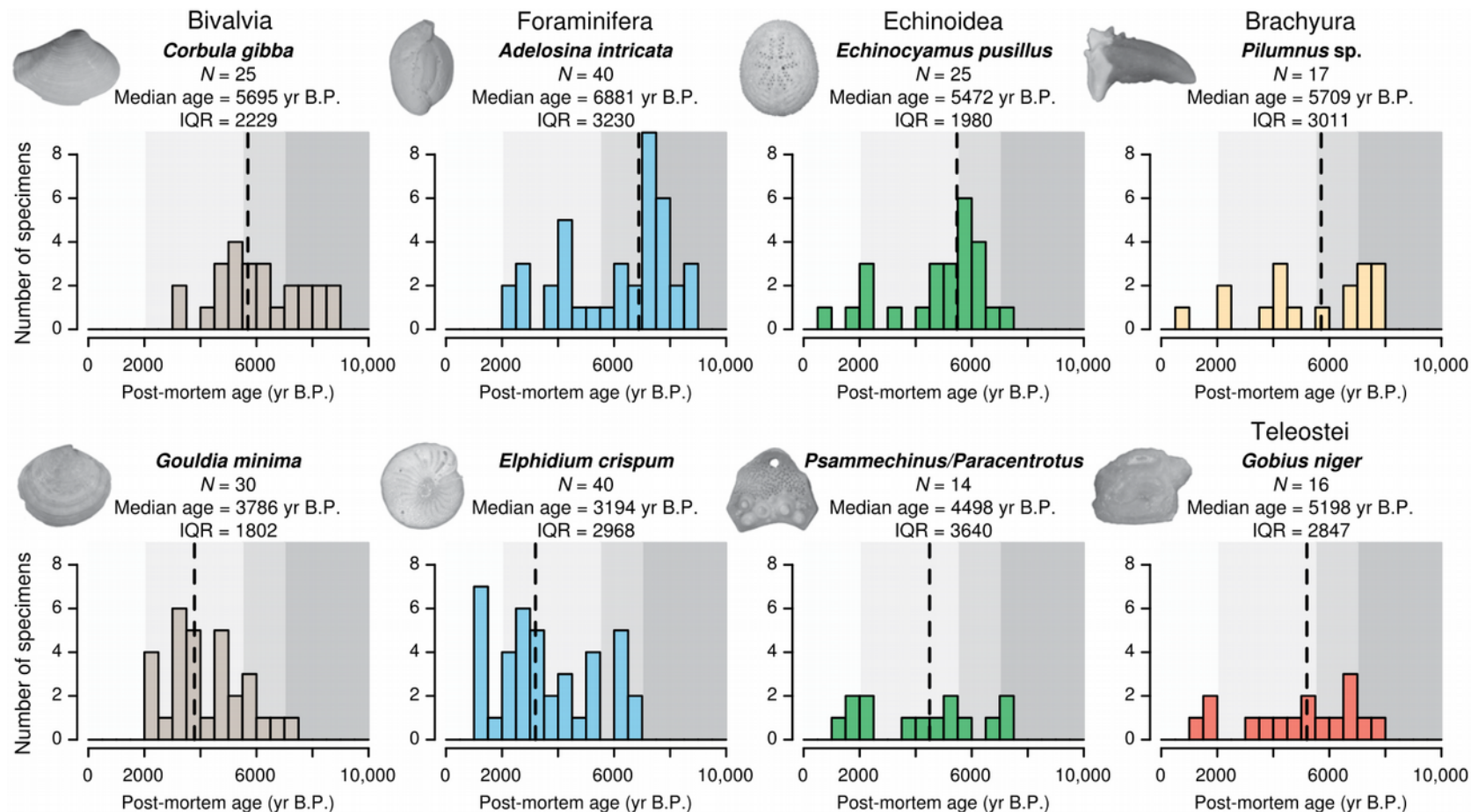


# Results: significant age offsets

Sea-level  
phases



Median ages differ between the taxa by up to ~3700 yrs



## Summary: similar temporal resolution but different average age

Within-taxon time averaging (IQR)

min: 1800 yrs → bivalve shells

max: 3640 yrs → regular echinoid plates

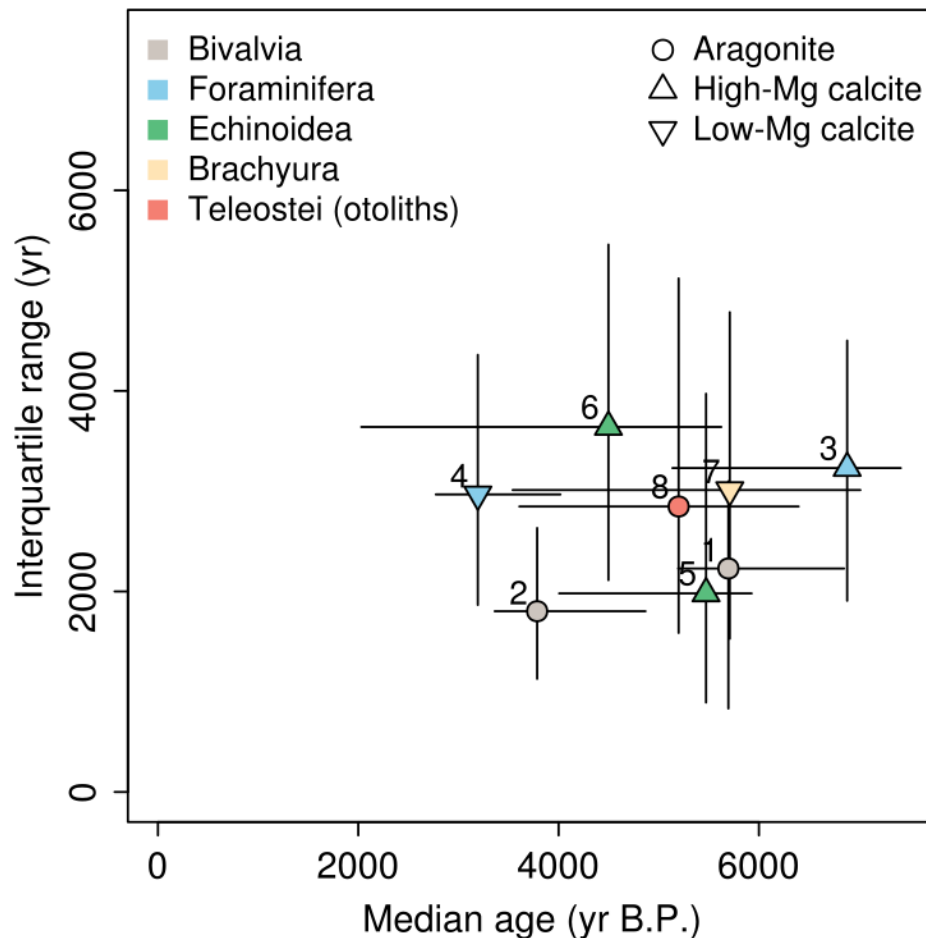
Age offset (difference between median ages)

min: 14 yrs → between the crab and bivalve

max: 3700 yrs → between the two forams

No relationship with size of the remains  
or skeletal mineralogy

Median age vs. IQR  $\pm$  95% bootstrapped CIs

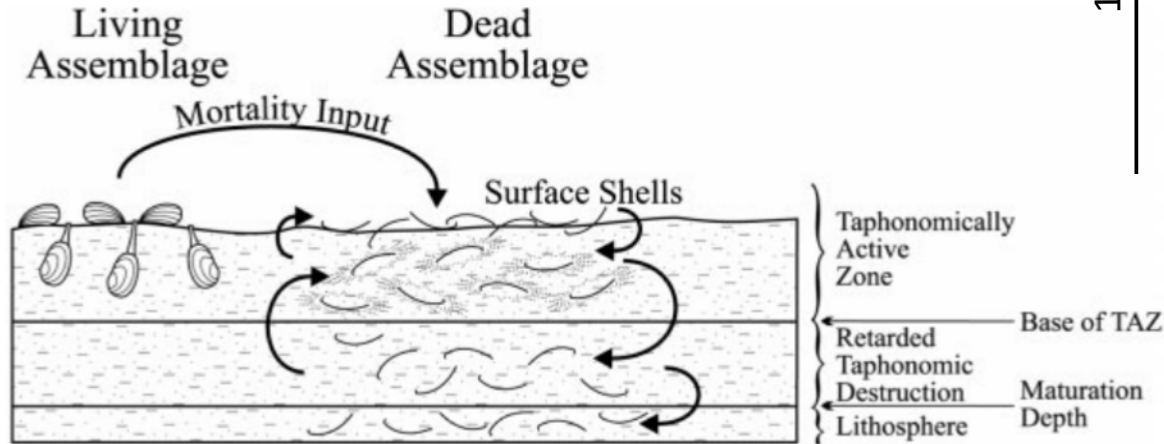




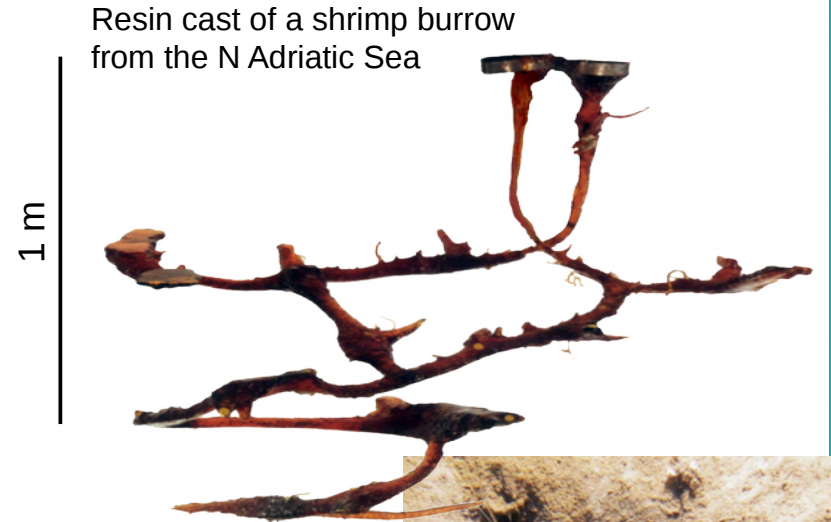
# Controlling factors: sequestration

Sequestration decouples time averaging from the effects of skeletal durability:

- **burial or early diagenetic processes** in the subsurface **protects shells** from rapid destruction in the taphonomic active zone
- older shells are occasionally **exhumed and mixed with younger cohorts** by deep bioturbation



Olszewski 1999 *Paleobiology*



Pervesler & Dworschak 1985  
*Senckenbergiana marit.*

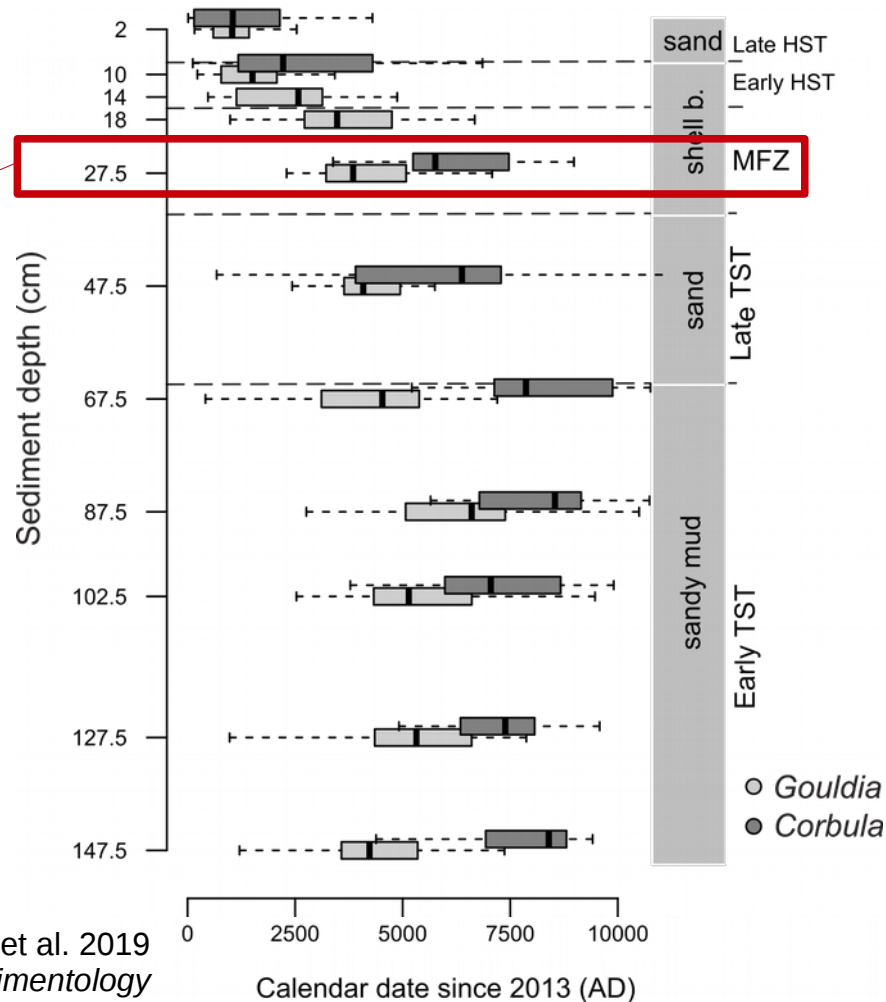
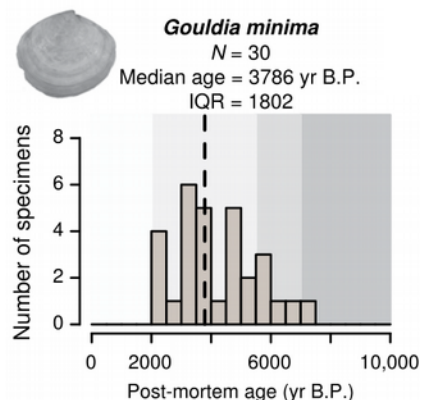
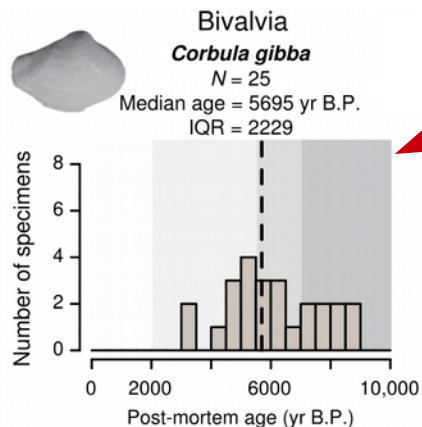




# Controlling factors: variability in skeletal production

Large **age offsets** between the two bivalve species are present throughout the studied core

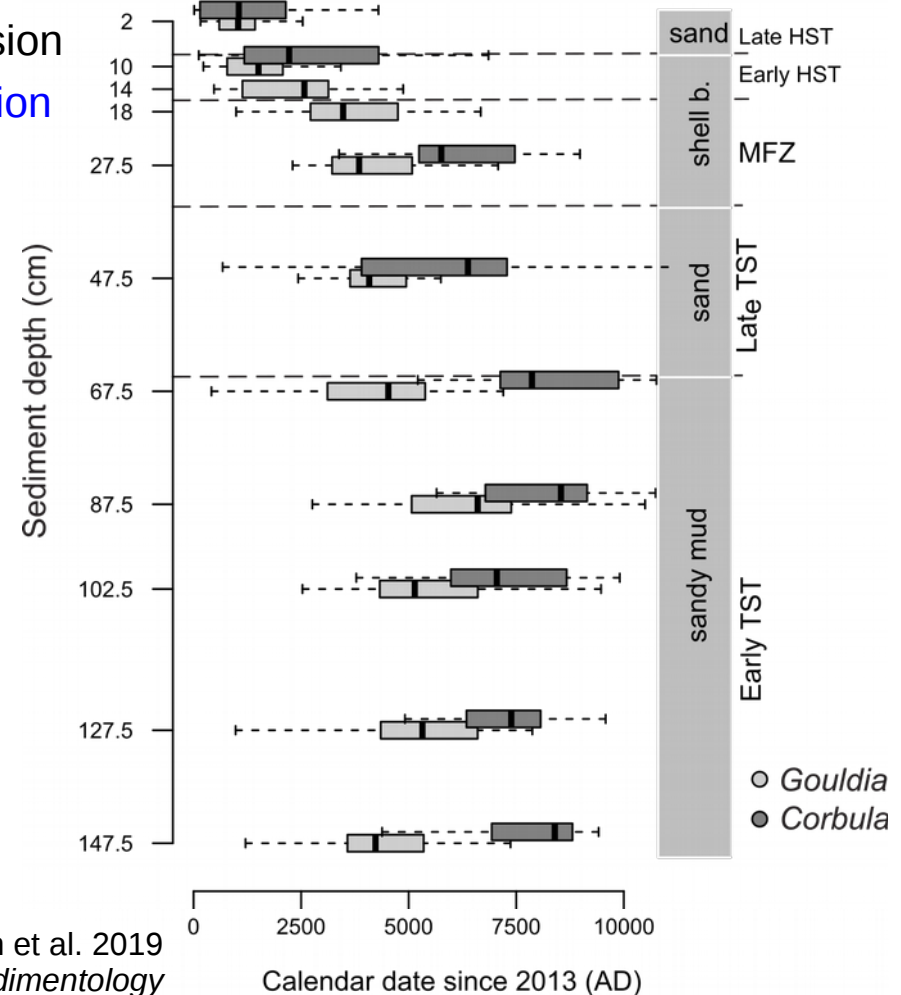
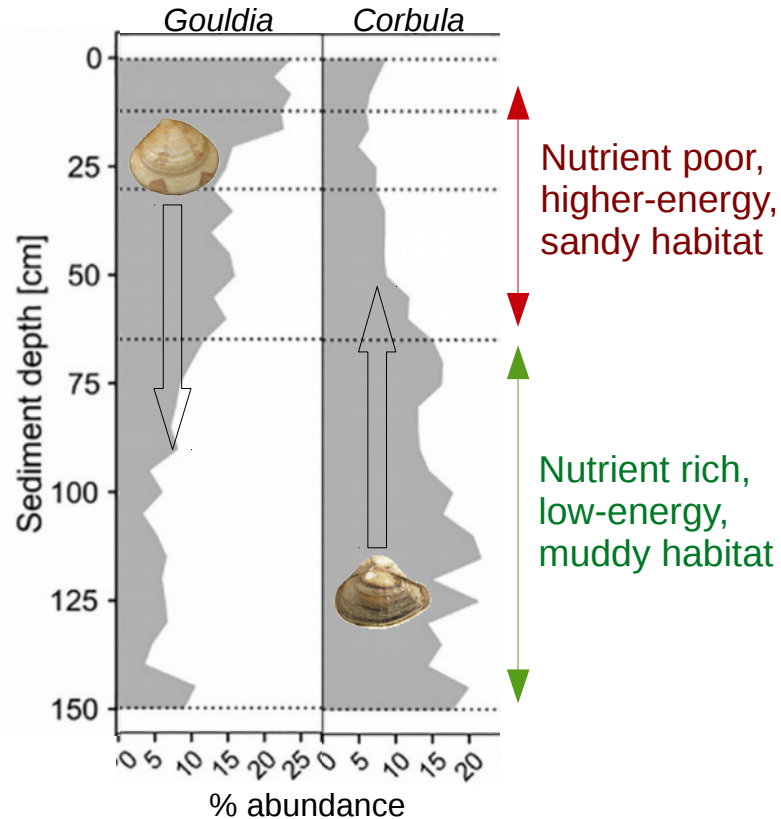
→ effect of **diachronous shifts in abundance**



Tomašových et al. 2019  
*Sedimentology*

# Controlling factors: variability in skeletal production

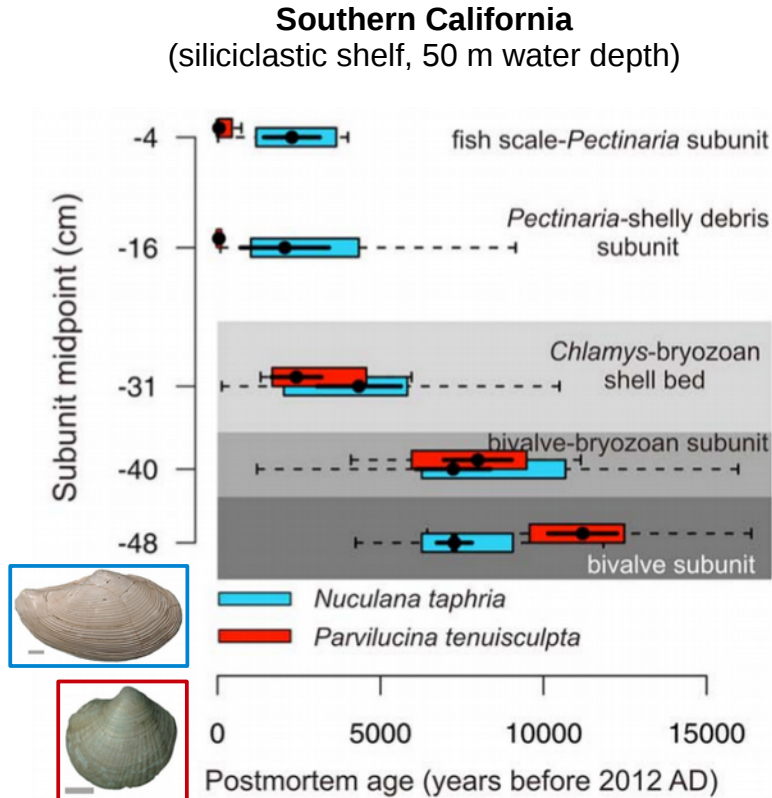
- peaks of abundance (and thus shell production) occurred at different times during the transgression
- shells of different ages mixed by deep bioturbation



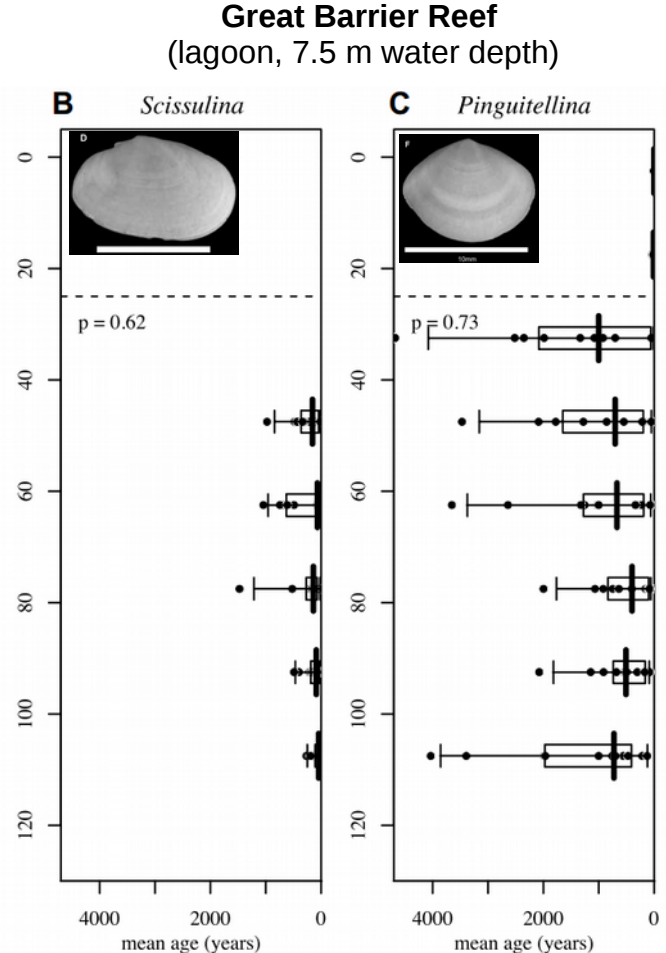
Tomašových et al. 2019  
*Sedimentology*

# Adriatic shelf is not unique

Age differences between co-occurring taxa are commonly observed in Holocene assemblages on continental shelves



Tomašových et al. 2019 *Paleoceanogr Paleoclimatol*



Kosnik et al. 2013 *Quat Geochronol*

# Conclusions



## Bad news

Just like macrofossils, **microfossil assemblages** can be **time-averaged over centuries to millennia** in shallow-water environments

**Significant age offsets between** co-occurring species can:

- complicate paleoecological and paleoenvironmental inferences
- make **age models** and **geochemical proxy** records **sensitive to taxon choice**

In other words, high-resolution age models based on dating of a single taxon may not be directly applicable to other members of the fossil assemblage or to geochemical proxy records based on their biomineralized hardparts

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## Good news

Fossil records of taxa with fundamentally different skeletal architecture and durability can have **comparable temporal resolution**

Limitations imposed by time averaging can be circumvented by direct **age dating of large numbers of specimens**

→ now possible thanks to recent advances in radiocarbon methods