

 Use independent tracers to validate chronology

- Check if depth distribution of other elements
- is homogeneous with depth.

  ✓ Analyze short-lived radionuclides at the SMI
- ✓ Check for burrowing evidence → e.g. X-ray radiographs or visual description of sed.

  core
- ✓ CRS or CF:CS model below SML → check residence time of <sup>210</sup>Pb in SML.
- ✓ The profile is likely undatable if entirely affected by mixing.

*(*4)

- Analyze short-lived radionuclides.
- ✓ Analyze <sup>226</sup>Ra if supported <sup>210</sup>Pb is not reached.
- Check historical records of event sedimentation.
- The profile is likely undatable by <sup>210</sup>Pb if other chronological tools are unavailable.

- (5)
- Check if changes in grain size distribution,
   DBD and OM content also occur.
- ✓ Normalize radionuclide concentrations to the parameter driving <sup>210</sup>Pb<sub>xs</sub> distribution.
- ✓ Analyze <sup>226</sup>Ra in all sections.
- ✓ CF:CS model in normalized profiles.
- ✓ The profile is likely undatable if normalization fails and other chronological tools are unavailable.

(7)

- Compare <sup>210</sup>Pb<sub>xs</sub> inventories with those at a reference site.
- Check for coarser grain size.
- Presence of short-lived radionuclides at reference site to check entire core recovery.
- ✓ CF:CS or CIC models to estimate mean MAR.

- (3)
- Use independent tracers to validate MAR in regions with different slope.
- Check for variations in geochemical or physical parameters in the profile.
- ✓ Test sensitivity to acceleration according to DBD and apparent MAR.
- ✓ Check historical records of natural or anthropogenic events.
- ✓ Apply CRS or CF:CS model piecewise.

(6)

✓ Only if OM is high (> 30 %) and mostly labile (0.01–0.03 d<sup>-1</sup>)

(8

- ✓ Analyze radionuclides in the fine sediment fraction (sieve to <63 μm or <125 μm) → check new profile
- √ <sup>226</sup>Ra by gamma spectroscopy
- ✓ Check for records of event sedimentation
- ✓ If sieving increases <sup>210</sup>Pb<sub>xs</sub> concentrations but its shape is similar, the profile is undatable and cannot be used for accumulation rate calculations.