

EDIT - feature request #6712

[DISCUSS] Should we remove anamorph from TaxonName (Fungi)

06/09/2017 01:44 PM - Andreas Müller

Status:	New	Start date:	
Priority:	New	Due date:	
Assignee:	Andreas Müller	% Done:	0%
Category:	cdm	Estimated time:	0:00 hour
Target version:	CDM UML 5.43		
Severity:	normal		
Description			
According to Melbourne Code anamorph should not be used anymore, or do we need it for older data.			
"One fungus, one name" and "one fossil, one name" are important changes for fungi and for fossils; the concepts of anamorph and teleomorph (for fungi) as well as morphotaxa (for fossils) have been eliminated.			
https://en.wikipedia.org/wiki/International_Code_of_Nomenclature_for_algae_fungi_and_plants			
Related issues:			
Related to EDIT - bug #6297: [Overview] BGBM User group wishes 2017			New

History

#1 - 06/09/2017 01:45 PM - Andreas Müller

- Related to bug #6297: [Overview] BGBM User group wishes 2017 added

#2 - 05/08/2018 06:55 PM - Andreas Müller

- Target version changed from CDM UML 5.0 to CDM UML 5.5

#3 - 05/05/2019 11:18 AM - Andreas Müller

- Target version changed from CDM UML 5.5 to CDM UML 5.15

#4 - 06/23/2020 02:53 PM - Andreas Müller

- Target version changed from CDM UML 5.15 to CDM UML 5.43