

## EDIT - Issues

| #     | Tracker         | Status   | Priority   | Subject   | Assignee         | Target version              | Category   | Tags                               |
|-------|-----------------|----------|------------|---|------------------|-----------------------------|------------|------------------------------------|
| 6166  | feature request | Closed   | Highest    | In ABCD import also import all siblings of a culture or tissue sample   | Katja Luther     | Release 4.7                 | cdmadapter | algoterra,abcd                     |
| 6399  | feature request | Closed   | Priority14 | Add admin property frequently used biocase provider   | Katja Luther     | Release 4.7                 | taxeditor  | abcd,gbol                          |
| 7433  | feature request | Resolved | Highest    | new service method to get specimen by genetic AccessionNumber   | Andreas Müller   | Release 5.8                 | cdmlib     | gbol,abcd                          |
| 7019  | feature request | Closed   | Priority14 | sort import menu  | Patrick Plitzner | Release 5.11                | taxeditor  | abcd                               |
| 10393 | bug             | Closed   | Highest    | ABCD Import of specimen added to a taxon included in second classification is not the one of the chosen classification        | Katja Luther     | Release 5.41                | cdmadapter | abcd,specimen                      |
| 10398 | feature request | Closed   | New        | Make it configurable in ABCD import if identifications should be imported as determinations or individualAssociations or both | Katja Luther     | Release 5.41                | cdmadapter | abcd,specimen                      |
| 8692  | bug             | New      | New        | Specimen online import is slow  | Katja Luther     | Unassigned CDM tickets      | taxeditor  | import,abcd                        |
| 6079  | bug             | New      | Priority14 | [Specimen Search] Restructure the table of search result  | Katja Luther     | Unassigned CDM tickets      | taxeditor  | abcd,import,specimen               |
| 10397 | bug             | Closed   | Highest    | Moving a specimen imported via ABCD to a new taxon does not show the specimen on the new taxon page but on the old one        | Katja Luther     | Release 5.42                | cdmadapter | specimen,abcd,specimens-view-table |
| 6325  | bug             | New      | Priority14 | Handling of specimen imported via biocase with missing nomenclatural code   | Katja Luther     | Reviewed Next Major Release | cdmadapter | abcd,gbol                          |
| 7434  | feature request | New      | Highest    | while importing sequences it should be possible to download fasta file  | Katja Luther     | Reviewed Next Major Release | cdmadapter | abcd,gbol                          |