

EDIT - Issues

| # | Tracker | Status | Priority | Subject | Assignee | Target version | Category | Tags |
|-------|-----------------|----------|------------|---|------------------|-----------------------------|------------|------------------------------------|
| 7433 | feature request | Resolved | Highest | new service method to get specimen by genetic AccessionNumber | Andreas Müller | Release 5.8 | cdmlib | gbol,abcd |
| 8692 | bug | New | New | Specimen online import is slow | Katja Luther | Unassigned CDM tickets | taxeditor | import,abcd |
| 6325 | bug | New | Priority14 | Handling of specimen imported via biocase with missing nomenclatural code | Katja Luther | Reviewed Next Major Release | cdmadapter | abcd,gbol |
| 6079 | bug | New | Priority14 | [Specimen Search] Restructure the table of search result | Katja Luther | Unassigned CDM tickets | taxeditor | abcd,import,specimen |
| 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 |
| 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 |
| 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 |
| 7019 | feature request | Closed | Priority14 | sort import menu | Patrick Plitzner | Release 5.11 | taxeditor | abcd |
| 6399 | feature request | Closed | Priority14 | Add admin property frequently used biocase provider | Katja Luther | Release 4.7 | taxeditor | abcd,gbol |
| 6166 | feature request | Closed | Highest | In ABCD import also import all siblings of a culture or tissue sample | Katja Luther | Release 4.7 | cdmadapter | algaterra,abcd |