

EDIT - Issues

| # | Tracker | Status | Priority | Subject | Assignee | Target version | Category | Tags |
|------|-----------------|-------------|------------|---|---------------------|-----------------------------|----------------|-----------|
| 6325 | bug | New | Priority14 | Handling of specimen imported via biocase with missing nomenclatural code | Katja Luther | Reviewed Next Major Release | cdmadapter | abcd,gbol |
| 6399 | feature request | Closed | Priority14 | Add admin property frequently used biocase provider | Katja Luther | Release 4.7 | taxeditor | abcd,gbol |
| 8607 | task | New | New | Dataprocessing for GBOL Dataportal/Algaterra | Wolf-Henning Kusber | Unassigned CDM tickets | data | gbol |
| 8182 | feature request | In Progress | Highest | show blast search result | Katja Luther | Release 5.45 | cdm-dataportal | 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 |
| 7433 | feature request | Resolved | Highest | new service method to get specimen by genetic AccessionNumber | Andreas Müller | Release 5.8 | cdmlib | gbol,abcd |