

EDIT - Issues

#	Tracker	Status	Priority	Subject	Assignee	Target version	Category	Tags
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
6325	bug	New	Priority14	Handling of specimen imported via biocase with missing nomenclatural code	Katja Luther	Reviewed Next Major Release	cdmadapter	abcd,gbol
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
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
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
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
6079	bug	New	Priority14	[Specimen Search] Restructure the table of search result	Katja Luther	Unassigned CDM tickets	taxeditor	abcd,import,specimen
7019	feature request	Closed	Priority14	sort import menu	Patrick Piltzner	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