

## 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
7433	feature request	In Progress	Highest	new service method to get specimen by genetic AccessionNumber	Katja Luther	Release 5.7	cdmlib	gbol,abcd
7019	feature request	New	Priority14	sort import menu	Katja Luther	Unassigned CDM tickets	taxeditor	abcd
6399	feature request	Closed	Priority14	Add admin property frequently used biocase provider	Katja Luther	Release 4.7	taxeditor	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
6166	feature request	Closed	Highest	In ABCD import also import all siblings of a culture or tissue sample	Katja Luther	Release 4.7	cdmadapter	alгатerra,abcd