

D4Science Infrastructure - Incident #6061

Cannot save and alter a shared Workspace folder

Nov 28, 2016 11:26 AM - Gianpaolo Coro

Status:	Closed	Start date:	Nov 28, 2016
Priority:	Urgent	Due date:	
Assignee:	Gianpaolo Coro	% Done:	100%
Category:	Data Management	Estimated time:	0.00 hour
Target version:	UnSprintable		
Infrastructure:	Production		

Description

The Home / SAI Examples / Simple_WPS4R_Annotation folder is shared among several people with "Write Any" rights. However, when opening this folder with SAI, a user who is not the owner of the folder gets a message like

"Insufficient Privileges to READ the node
/Home/gianpaolo.coro/Workspace/StatisticalAlgorithmsImporter/Simple_WPS4R_Annotation/AbsencesSpeciesList-prod-annotated.r
for user ..."

whereas, the owner of the folder cannot alter the files in the folder. When overwriting, no error is returned but the file is not changed.

History

#2 - Nov 28, 2016 11:28 AM - Gianpaolo Coro

This issue is similar to [#5626](#), which was solved by restarting the portal.

#3 - Nov 28, 2016 11:57 AM - Pasquale Pagano

- Priority changed from Normal to Urgent

#4 - Nov 28, 2016 12:05 PM - Valentina Marioli

- Status changed from New to In Progress

#5 - Nov 28, 2016 04:36 PM - Valentina Marioli

- Status changed from In Progress to Feedback

- Assignee changed from Valentina Marioli to Gianpaolo Coro

- % Done changed from 0 to 100

"SAI Examples/Simple_WPS4R_Annotation" is now working.

The problem wasn't related to HL, but to wrong references to the project in the file "SAI Examples/Simple_WPS4R_Annotation/stat_algo.project". Except for the root folder "SAI Examples/Simple_WPS4R_Annotation", the other references were to a private folder of gianpaolo.coro "/Workspace/StatisticalAlgorithmsImporter/" (paths and ids). Such folder is not a shared folder, so it's right that you get an Insufficient Privileges Exception for every user except "gianpaolo.coro", the owner.

Maybe the issue is related to the bug opening multiple tabs. Such bug will be fixed in gCube 4.2.

I've fixed the problem modifying the file "SAI Examples/Simple_WPS4R_Annotation/stat_algo.project", replacing the wrong paths and ids with the correct files in "SAI Examples/Simple_WPS4R_Annotation".

#6 - Nov 28, 2016 04:54 PM - Gianpaolo Coro

- Status changed from Feedback to In Progress

Saving the main code to the WS works, but the recompilation does not affect the process executed by DataMiner. The package in the Target/Deploy folder is changed (I have checked the content), but this does not correspond to the file downloaded by DataMiner, which is

<http://data.d4science.org/bksxMGVWTI3WjM5Z1ZXYXUOUtMZzVSNIBhZXFGb05HbWJQNStIS0N6Yz0>

Indeed, the files should be exactly the same. What file and folder the link above is really pointing to?

#7 - Nov 28, 2016 04:54 PM - Gianpaolo Coro

- Assignee changed from Gianpaolo Coro to Valentina Marioli

#8 - Nov 28, 2016 05:30 PM - Valentina Marioli

<http://data.d4science.org/bksxMGVWTIZ3WjM5Z1ZXYXIUOUtMZzVSNIBhZXFGb05HbWJQNSiIS0N6Yz0> is a link to
/Home/gianpaolo.coro/Workspace/StatisticalAlgorithmsImporter/Simple_WPS4R_Annotation/Target/Deploy/AbsencegenerationfromOBIS.zip .

#9 - Nov 28, 2016 06:04 PM - Valentina Marioli

- Status changed from *In Progress* to *Feedback*
- Assignee changed from *Valentina Marioli* to *Gianpaolo Coro*

#10 - Nov 28, 2016 06:19 PM - Gianpaolo Coro

- Status changed from *Feedback* to *Resolved*

You are right, that was my mistake! Thank you very much.

#11 - Nov 28, 2016 06:21 PM - Valentina Marioli

- Status changed from *Resolved* to *Closed*