

Kepler - Bug #2253

ENM Workflows errors on newer, faster computer

11/07/2005 09:32 AM - Dan Higgins

<b>Status:</b>	Resolved	<b>Start date:</b>	11/07/2005
<b>Priority:</b>	Normal	<b>Due date:</b>	
<b>Assignee:</b>	Dan Higgins	<b>% Done:</b>	0%
<b>Category:</b>	general	<b>Estimated time:</b>	0.00 hour
<b>Target version:</b>	1.0.0alpha8	<b>Spent time:</b>	0.00 hour
<b>Bugzilla-Id:</b>	2253		
<b>Description</b> Errors are occurring in 'SingleSpecies_BestRuleSet_Aug2005 ENM workflow on new, faster computer (Optiiplex GX620 - 3GHz dual core - Win XP) that do not occur when same workflow is run on older, slower machine (null pointer error most of time). Error seems to occur at end of GARP run; perhaps a file is not ready?  Possible threading problem?			

History

#1 - 11/08/2005 12:42 PM - Dan Higgins

Found the problem in the RExpression actor (after 2 1/2 days of looking!). Needed to add another \_read statement before looking at the output string from the R process. Apparently, some text was added in time between read thread and main Kepler thread which resulted in mission output from RExpression actor.

#2 - 03/27/2013 02:19 PM - Redmine Admin

Original Bugzilla ID was 2253