

Kepler - Bug #2139

mollusc -> R -> plot WF for examples and documentation uses

07/05/2005 02:17 PM - Chad Berkley

Status:	Resolved	Start date:	07/05/2005
Priority:	Normal	Due date:	
Assignee:	Samantha Romanello	% Done:	0%
Category:	documentation	Estimated time:	0.00 hour
Target version:	1.0.0rc1	Spent time:	0.00 hour
Bugzilla-Id:	2139		
Description			
We need a simple R workflow for use in examples and documentation. This may already be built and simply needs to be updated for ease of use.			

History

#1 - 11/02/2005 05:12 PM - Matt Jones

The mollus workflow that uses GCE mollusc data and then plots it in R could be expanded to be a bit more realistic, and documented well for the tutorials.

#2 - 06/14/2006 11:02 AM - Dan Higgins

There is now documentation for using the RExpression actor (in Kepler-docs/user/R_Actor_userdoc_draft.doc) that uses the GCE data, but from a local file rather than the ecogrid (due to problems with the ecogrid version which have now been fixed). Should be updated.

#3 - 04/06/2007 01:55 PM - Dan Higgins

R workflow is now in the Getting Started Guide

This specific bug seems fixed.

#4 - 03/27/2013 02:19 PM - Redmine Admin

Original Bugzilla ID was 2139