

SEEK - Bug #1068

create and analyze example pipelines with workflow editor

05/20/2003 10:00 AM - Matt Jones

<b>Status:</b>	Resolved	<b>Start date:</b>	05/20/2003
<b>Priority:</b>	Normal	<b>Due date:</b>	
<b>Assignee:</b>	Deana Pennington	<b>% Done:</b>	0%
<b>Category:</b>	ams	<b>Estimated time:</b>	0.00 hour
<b>Target version:</b>	Unspecified	<b>Spent time:</b>	0.00 hour
<b>Bugzilla-Id:</b>	1068		
<b>Description</b> Need to create several representative formal workflows using the workflow editor described in bug 1064 to show what typical workflow scenarios are. One or two of these should be focused around the GARP case study. Several others should be from other areas of ecology to get some breadth.  Once these have been created, need to: 1) review workflows to see if scientists agree that the representation is correct 2) analyze the workflows to see if there are constructs that were difficult or impossible (indicating the need for a more expressive system) 3) analyze the workflows for use in developing semantic type ontologies			
<b>Related issues:</b> Blocks SEEK - Bug #1064: create initial workflow editor from Vergil			
		<b>Resolved</b>	<b>05/20/2003</b>

History

#1 - 02/12/2004 08:16 AM - Deana Pennington

Abstract workflow for mammal project completed at BEAM workshop, 2/6/04.

#2 - 03/27/2013 02:16 PM - Redmine Admin

Original Bugzilla ID was 1068