

Kepler - Bug #2252

Need to add actors for aligning multiple DNA or protein sequences

11/06/2005 06:33 PM - Timothy McPhillips

Status:	Resolved	Start date:	11/06/2005
Priority:	Normal	Due date:	
Assignee:	Timothy McPhillips	% Done:	0%
Category:	actors	Estimated time:	0.00 hour
Target version:	Post-rel-1.0.0	Spent time:	0.00 hour
Bugzilla-Id:	2252		
Description			
We need to wrap the ClustalW and Gblocks programs in collection-oriented actors. The aligned sequences output by these actors could then be fed into the actors wrapping the Phylip package and used for inferring phylogenetic trees.			

History

#1 - 10/23/2008 04:46 PM - Timothy McPhillips

Actors for running ClustalW and Gblocks are included in the ppod-actors module.

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

Original Bugzilla ID was 2252