



**Integrative Cancer Research Special Interest Group Teleconference**

**Proteomics SIG Meeting Minutes**

<b>Date, Time &amp; Location:</b>	September 13, 2004 2:00 – 3:00 PM EDT																		
<b>Attendees:</b>	Patrick McConnell – Duke Simon Lin – Duke Jennifer Shoemaker – Duke Michael Ochs – Fox Chase Jason Grundstead – U of Iowa Holden Juli Klemm - BAH																		
<b>Updates from last meeting</b>	<ul style="list-style-type: none"> <li>- Simon Lin met with representatives from the National Institute for Environmental Health. They have developed an object model to support systems biology research. They have combined elements from the MIAME and MIAPE standards, and are interested in adding in Duke's SMOS model as a plug-in to theirs. They have prepared a paper that has been accepted for publication in Bioinformatics. The citation is posted on the caBIG Proteomics SIG discussion forum: <a href="http://ncicbforums.nci.nih.gov/cabigforum/fcForum/0663978480991/407588140618/view_html?forum_loc=0663978480991">http://ncicbforums.nci.nih.gov/cabigforum/fcForum/0663978480991/407588140618/view_html?forum_loc=0663978480991</a></li> </ul>																		
<b>Future Directions</b>	<ul style="list-style-type: none"> <li>- The group discussed adoption of the MIAPE model interactions between the RProteomics project and the Proteomics LIMS project.</li> <li>- The LIMS project will likely leverage the efforts of the RProteomics group at this time from an intellectual perspective. They will eventually adopt the MIAPE model, but probably not in the next 6 months. With regard to common data elements, it may still be early too try to agree on these.</li> <li>- LIMS group is starting from the caLIMS data schema, but may need to modify it.</li> <li>- The group agreed that caCORE/caBIO training would be of great benefit.</li> <li>- Patrick McConnell and Michael Ochs agreed that seeing each other's data models in the near term would be beneficial and it was agreed they would both send their current models to all SIG members.</li> <li>- Patrick McConnell pointed out that it would be a good idea for this group to start thinking about use cases related to interoperability of projects within this SIG. The group agreed to craft preliminary use cases of this sort for the use cases due to the Architecture group this Friday, 9/17/04.</li> </ul>																		
<b>Proteomics SIG lead</b>	<ul style="list-style-type: none"> <li>- Michael Ochs volunteered to Fox Chase be in charge of the Proteomics SIG meetings moving forward. He will identify an individual from his team to play this role. They will generally be responsible for organizing and facilitating the meeting and creating meeting minutes.</li> </ul>																		
<b>Action Items</b>	<table border="1"> <thead> <tr> <th>Name Responsible</th> <th>Action Item</th> <th>Date Due</th> <th>Notes</th> </tr> </thead> <tbody> <tr> <td>Juli Klemm</td> <td>Distribute meeting minutes</td> <td>9/20/04</td> <td></td> </tr> <tr> <td>Michael Ochs, Patrick McConnell</td> <td>Distribute current data schemas to all members of the Proteomics SIG</td> <td>9/24/04</td> <td></td> </tr> <tr> <td>Shannon McWeeney</td> <td>Summarize all proteomics data that would be available for</td> <td>Next SIG meeting</td> <td></td> </tr> </tbody> </table>			Name Responsible	Action Item	Date Due	Notes	Juli Klemm	Distribute meeting minutes	9/20/04		Michael Ochs, Patrick McConnell	Distribute current data schemas to all members of the Proteomics SIG	9/24/04		Shannon McWeeney	Summarize all proteomics data that would be available for	Next SIG meeting	
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		sharing from Oregon Health			
	Michael Ochs	Identify an individual from Fox Chase who will be in charge of the Proteomics SIG meetings moving forward	9/20/04		
	All SIG members	Include ideas for how projects in this SIG might interoperate when putting together use cases to share with the Architecture group	9/17/04		