Open Science Grid All Hands Timetable

Tuesday, 9 March 2010

<u>Biology Applications</u> - Wilson Hall, One North - North West (08:30 - 12:00)

time	[id] title	presenter
08:30	[16] Welcome and Goals	MCGEE, John SLIZ, Piotr
08:40	[17] Practical approach to privacy, access, and processing for biological synchrotron experiments.	BRISTER, Keith
09:00	[18] Shotgun Metagenomics - TBD	MEYER, Folker
09:20	[19] Production Computational Biology Workflow on OSG; Experience, Tools and Perspectives	STOKES-REES, Ian
09:40	[20] Genome Center at WUSTL	STIEHR, Gary
09:50	[58] Molecular basis for uninducible behavior of mutations at position 15 in AraC protein	MILLER, Tim
10:00	Break	
10:15	[21] Discriminative Learning for Protein Conformation Sampling	XU, Jinbo
10:35	[47] DIAG: A data intensive academic grid - A platform for performing bioinformatics analysis on genome sequence data	MAHURKAR, Anup
10:55	[23] MPI Cluster for Molecular Dynamics Computations	DOHERTY, Peter
11:05	[22] Desmond: An Efficient, Scalable Code for Biomolecular Simulation - KEYNOTE	BOWERS, Kevin
11:50	[24] Discuss the value a collaboration with OSG has and/or can bring to these computing needs.	