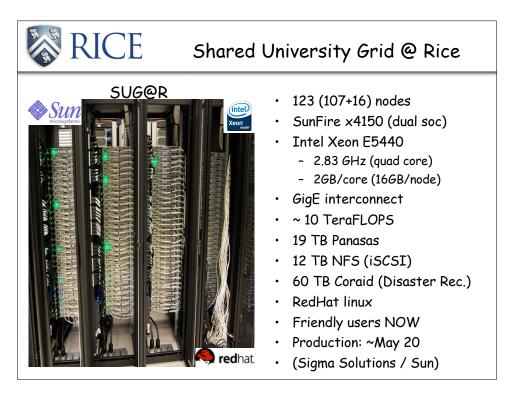
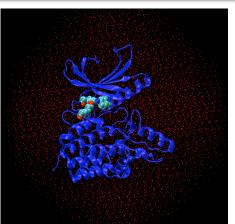


RICE "HPC" Resources (12 months ago)						
Manufacturer	Architecture	os	CPU	Memory (GB)	Local Disk (TB)	% tot
Cray	AMD Opteron	SuSE	672	1400	26	45%
HP	Intel Itanium	RedHat	278	608	8	19%
Linux Networx	Intel Xeon	RedHat	84	84	1.2	6%
Appro	Intel Xeon	RedHat	150	300	1	10%
Appro	AMD Athlon	RedHat	84	256	1	6%
Appro	AMD Athlon	RedHat	44	44	0.5	3%
Appro	AMD Opteron	RedHat	32	64	4	29
PSSC	Intel Xeon	RedHat	24	48	1.5	29
Appro	Intel Xeon	RedHat	24	48	1.5	2%
Appro	AMD Athlon	RedHat	20	40	1	17
PSSC	AMD Opteron	Fedora	20	40	0.5	19
IBM	Regatta	AIX	16	32	1	17
IBM	Regatta	AIX	16	64	1	17
Chaogic	AMD Athlon	RedHat	8	8	0.2	17
IBM	Power5	AIX	4	8	1	0%
Dell	PowerEdge 6850	Windows	4	16	0.5	0%
Total for all Compute Clusters			1480	3060	56.1	100%



## RICE Molecular Dynamics of Proteins

- Compute type and strength of interactions between kinase and drug through time (20-40ns)
- Calculate binding free energy of the drug from MD simulation.
- Amber 9
  - Pmemd: parallel version to do MD for systems with explicit water
- System (31908 atoms)
  - GAK kinase (4628 atoms)
  - inhibitor (55 atoms)



MD of G-associated **kinase** (GAK) in complex with inhibitor (Iressa) for treatment of lung cancer

