# Texas Advanced Computing Center (TACC) Overview

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Life Sciences Computing



## Texas Advanced Computing Center (TACC)

TACC's mission is to enable discoveries that advance science and society through the application of advanced computing technologies.

UT Austin: What starts here, changes the world

TACC: Powering discoveries that change the

world



#### TACC's Vision

- Provide the most powerful, capable computing technologies and techniques that enable people researchers, educators, developers, engineers, businessmen, etc.—to advance science and society.
- Provide leadership in the advanced computing community in technology R&D, support, education, and expertise to ensure maximum impact of current and future technologies in diverse applications.
- Enable transformational science and societal achievements that change, influence, and improve our understanding of the world, and the world itself.



#### **About TACC**

- Research Center within the University of Texas at Austin
- ~110 staff including ~30 PhD level researchers
- Manages computational resources for the scientific community and provides them <u>free of charge</u> to investigators on a project basis
- Conducts research on a variety of topics
- Funded by NSF, UT, other federal grants, and traditional PI research grants





# Ranger

- 62,976 cores, 2.0 GHz AMD Opteron
- Will retire in February 2013
- Still one of the most powerful supercomputers in academia, with over 579 teraflops peak performance.
- To build and deploy Ranger, the National Science Foundation awarded TACC \$59 million, the largest single award ever from that agency to UT Austin.



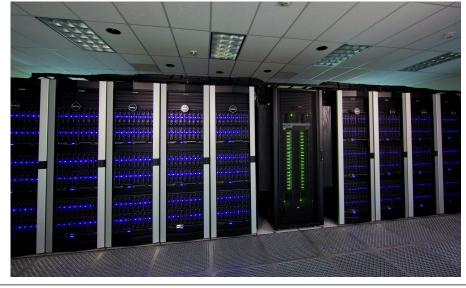
#### Four year statistics:

- 97% uptime
- Over 2 Billion jobs run
- More than 1.5 Billion hours of compute time used



#### Lonestar

- Partners include NSF, UT Austin, UT System, Texas A&M and Texas Tech
- 1,888 Dell M610 PowerEdge blade servers, each with two six-core Intel Xeon 5600 "Westmere" processors – total 22,656 cores
- Smaller than Ranger, but more powerful per node
- 302 teraflops peak performance





# Stampede

- Base Cluster (2 Petaflops):
  - Intel Sandy Bridge (8 core) processors
  - Dell dual-socket nodes w/32GB RAM
  - More than 6,000 nodes
  - More than 100,000 cores
- Co-Processors (8 Petaflops):
  - Intel Xeon Phi co-processors
  - Special release of "Knights Corner" (>50 cores)
- Total concurrency approaching 500,000 cores



# **UT Data Repository (Corral)**

- 5 Petabytes of geo-replicated storage
  - Data centers in Austin and in Arlington
  - Parallel file system on high-speed network
- Available to all MD Anderson researchers
- Allocations up to 5TB are free of charge
- Larger allocations for \$250/TB/year
- Synergistic with Lonestar/Stampede for compute and Ranch for archive storage



# **Data Storage Costs**

	TACC Corral	Google Drive	SkyDrive	Dropbox	iCloud
Free Basic					
Account	5TB	5GB	7GB	2GB	5GB
10GB					\$20
20GB			\$10		\$40
25GB		\$30			
50GB			\$25	\$100	\$100
100GB		\$60	\$50	\$200	\$200
1TB		\$600		\$795 (five users)	
2TB		\$1,200			
4TB	_	\$2,400		_	
5TB	FREE			_	
8TB	\$2,000	\$4,800			
16TB	\$4,000	\$9,600			



# Ranch Tape Archive

- 70 petabyte capacity (and growing)
- Used for long-term storage
- Access provided freely to users of other TACC
  - resources
- Data staged on spinning disk and stored long-term on tape





# O'Donnell Gift: Data-Driven Science

- Announced February 21, 2012
  - initial production Q4 2012
- \$10 M over three years for:
  - a high-performance petascale data storage system accessible from all of TACC's computing systems
  - a computing system with embedded high-speed storage, optimized for data-intensive computing
- \$2 M matching funds from UT for staffing



### Visualization on Longhorn

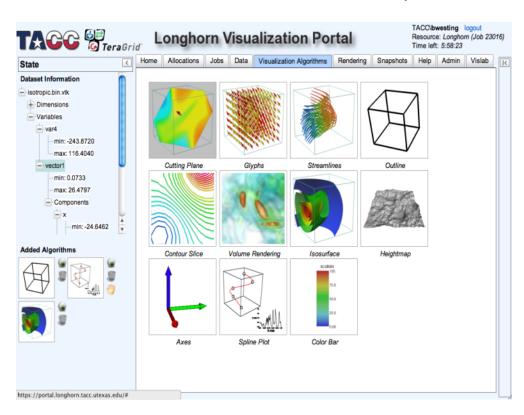
- The largest system in the world dedicated to visualization and data analysis
- System totals: 2048 cores, 13.5 TB distributed RAM, 512 GPUs
- 17M compute hours available annually
- Ideal for remote visualization and GPU computing



# Longhorn User Portal

#### Remote interactive visualization tool, developed at TACC

- Dramatically simplifies the visualization process
- Semi-automated data importation



- Powerful, easy-to-use entry point for researchers
- Quick and easy interactive visualization
- Web-based works from anywhere!



# Life Sciences Computing at TACC



#### www.tacc.utexas.edu/life-sciences-computing/applications

- TACC maintains a quarterly release cycle for a core set of biology related applications that:
  - Have high impact and broad appeal to life sciences research
  - Are highly requested by the user community
- Compiler optimized installations using fast math libraries when possible
- 80+ packages installed for genomics, phylogenetics, and computational chemistry



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#### **Genomics**

Abyss

**BEDTools** 

BioPerl

Bismark

Blat

Bowtie

**BWA** 

Cufflinks

**FASTX-Toolkit** 

**GATK** 

**GSNAP** 

**HMMER** 

Libgtextutils

**MAFFT** 

Maq

mpiBLAST

**MUSCLE** 

NCBI BLAST+

Newbler

**Oases** 

**Picard** 

SAMtools

**SHRIMP** 

SOAPdenovo

SRA toolkit

**SSAKE** 

**Tophat** 

TrinityRNASeq

Velvet



www.tacc.utexas.edu/life-sciences-computing/applications

#### **Phylogenetics**

BEAST RAxML-Light

**ClustalW** Parsimonator

MrBayes Phyutility

PHYLIP QuickTree

**RAxML** 



www.tacc.utexas.edu/life-sciences-computing/applications

#### **Computational Chemistry**

Amber GAMESS

APBS GROMACS

AutoDock LAMMPS

AutoDock Vina NAMD

Desmond NWChem

**Espresso** Siesta

Gaussian (coming...) VASP



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**Broadly Used Tools** 

idev

Java / JDK

launcher

Octave

R (statistics package)

...and many others



# User-Specific Software

- What if I need software not already installed?
  - Will it run on Linux?
  - Is the source code available?
  - Are there licensing restrictions?
- Try installing it in your personal directory (root access not required)
- Submit a consulting ticket on the TACC Portal for help
- Email or talk with one of us



# **Training Opportunities**

- Training provided at no cost for UT System researchers
- Several training classes per year in Houston
  - BYOCD Workshop 2/16/2012
- Web participation for training onsite at TACC
- Recordings are now frequently available at: <a href="http://www.tacc.utexas.edu/user-services/">http://www.tacc.utexas.edu/user-services/</a> <a href="training/course-materials">training/course-materials</a>



#### Research Collaboration

#### **Examples:**

- The iPlant Collaborative
- Epigenome Dynamics During DNA Replication
- Docking @ UTMB Web Portal
- RepServer
- National Center for Genome Analysis Support



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