

Genome Analysis

Birds of a feather

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National infrastructure serving genome science

Creators of new software

NCGAS – small, serving large community largely reactively

- Trinity
- Galaxy
- ABySS
- Velvet

iPlant – large collaborative serving plant science

- DNA Subway
- iPlant Discovery Environment
- · Many bioinformatics software applications planned as part of group strategy

XSEDE – designed to serve all research communities

- Stampede
- Gordon
- Blacklight
- Comet
- Mason
- Wrangler
- FutureGrid

Network – essentially independent of any particular research community

- Internet2
- · Regional providers



XSEDE resources

| System | Type of resource | Type of Service Provider? |
|------------|--|--------------------------------|
| Stampede | Large scale distributed memory parallel | NSF funded, Level 1 |
| Gordon | Large scale distributed memory parallel, pseudo large memory | " |
| Blacklight | Large memory | ш |
| Comet | New - VMs | u |
| Wrangler | Storage | u |
| FutureGrid | Experimental computer science / cloud system | и |
| Mason | Large memory (low cores) | IU-funded, Level 2 |
| Rockhopper | Commercial "cluster as a service" owned by Penguin Computing and housed at / supported by IU | Commercially owned, Level 3 |



Pittsburgh Supercomputing Center Blacklight (SGI Altix® UV 1000) - Massive Coherent Shared Memory Computer

2×16 TB of cache-coherent shared memory, 4096 cores

- ideal for genome sequence assembly
- High bandwidth, low latency interprocessor communication

SUSE Linux operating system

- excellent for portability:
 supports OpenMP, C, C++,
 Java, Perl, Python, p-threads,
 MPI, UPC
- rapid algorithm development



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Mason

 Supports data-intensive high performance computing tasks for IU researchers, faculty, staff, and students on all campuses.

Specs:

- Peak performance of 3.83 teraFLOPS
- 8 TB total RAM 512 GB RAM per node – really a system of memory with a few processors attached
- Uses Lustre/Data Capacitor II as high performance file system
- Connects to IU's high speed research network via 10 Gbps connection





iPlant – Plant Cyberinfrastructure

Goals:

- "to create a new type of organization a cyberinfrastructure collaborative for plant science"
- "to enable new conceptual advances through integrative, computational thinking"
- "to address an evolving array of grand challenge questions in plant science: the driving force and organizing principles for the collaborative"
- ~ \$10M / year (\$50M NSF Funded Project 5 years, renewed in 2013)
- iPlant is a cyberinfrastructure *platform*
- The platform is developed by iPlant and extensible by users
- NSF recommended scope beyond plants



XSEDE Novel and Innovative Projects program



- Novel and Innovative Projects within XSEDE is intended to be reactive to new user needs,
 with current focus on life sciences
- Work with developers to port key de novo assembly applications to large shared memory system, Blacklight
- Availability of Blacklight highlighted on Broad Institute developer web pages (ALLPATHS-LG and Trinity) and genomeweb.com
- Enthusiastic response from research community dozens of new groups using Blacklight for de novo assembly every year
- Example projects:
 - Cold Spring Harbor: Assembled 5 and 10 gigabase wheat species using 3 and 6 TB RAM respectively. Targeting assembly of 16 gigabase wheat genome (ALLPATHS-LG).
 - Cornell and Broad Institute: Assembled 20 primate transcriptomes at ~1 TB RAM each (Trinity). Understanding evolutionary processes and gaining insight into human disease.

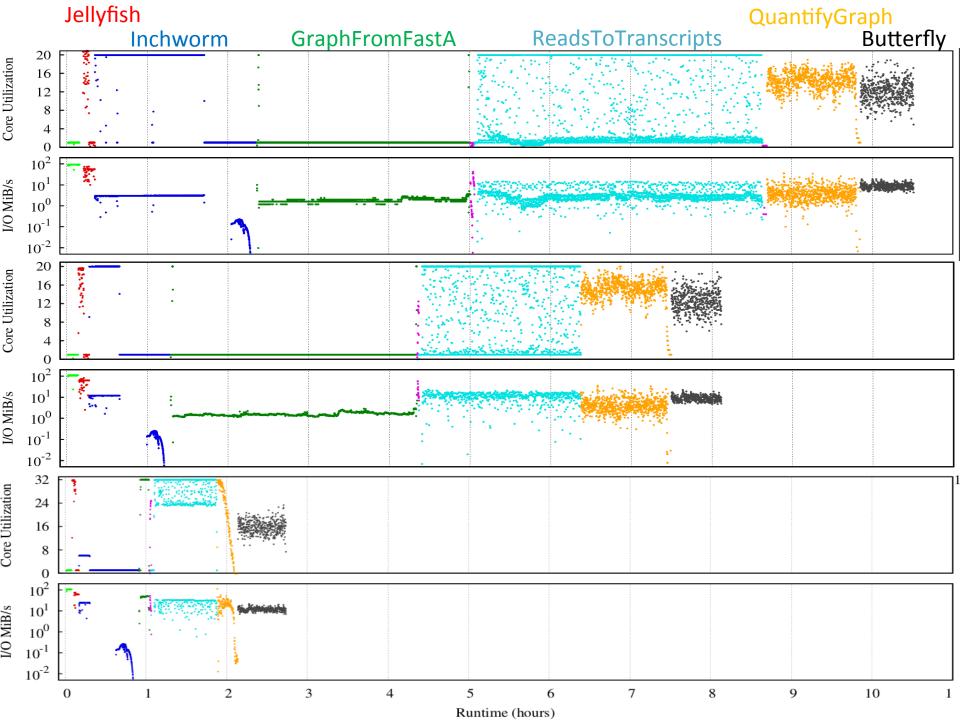
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National Center for Genome Analysis Support "Mind the Gap"

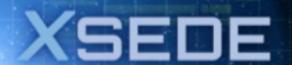
| Gap | How we fill it |
|---|---|
| System configurations offered by XSEDE and what people doing genome assembly need | Mason (IU contribution to facilities) |
| Software on XSEDE is not what people need | NCGAS installs and maintains |
| Software works slowly | NCGAS tunes / re-engineers |
| People just need help | NCGAS provides consulting NCGAS goes to conferences and informs people about our services |
| People need storage | NCGAS provides tape storage (IU facilities) |
| People need to publish data sets | IU provides resources via IUScholarWorks |





REUs

- NCGAS Virtual Interns leverage XSEDE experience to gain industry employment.
- NCGAS ran virtual REU program with Clark State University in Springfield OH, learning how to install and configure bioinformatics software.
- Two associate degree students participated.
- Both ended up working as professional HPC admins at Wright Patterson Air Force Base.



So... what are your pain points?

And what can XSEDE and NCGAS do to help?



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