



# GenomeHubs: Life Science Informatics & HPC

*HubBub 2013 Conference - <http://hubzero.org>*



A man with glasses and a black shirt is holding a camera up to his eye, taking a photo. He is standing in front of a wall of server racks. The text is overlaid on the left side of the image.

I m Chris."

"

I m an infrastructure geek."

"

I work for the BioTeam."

slideshare.net/chrisdag/ - [www.bioteam.net](http://www.bioteam.net) - Twitter: @chris\_da

# BioTeam

Who, What, Why ...

- ▶  **Independent consulting shop**
- ▶  **Staffed by scientists forced to learn IT, SW & HPC to get our own research done**
- ▶  **10+ years bridging the gap between science, IT & high performance computing**
- ▶  **We get to see how many groups of smart people tackle similar problems**







**DISCLAIMER.**



# Why I m here

Not a Hub Expert. Simply a Bio/IT Nerd

- ▶  **One of my clients decided to deploy HubZero last year**
  - I was involved in initial eval and source / VM based installs
- ▶  **Met HZ team virtually while:**
  - Hacking Submit framework to support the Grid Engine job scheduler
  - Building a [chef cookbook](#) to automate Hub installs onto Amazon EC2 node images
- ▶  **HubBub 13 folks invited me to talk life science HPC and where Hubs can play a role**
- ▶  **May also be a token Corporate/Industry/Enterprise science type**





# Goal: “Talk Fast & Get Out Of The Way”

## Quick Sprint

Life Science Informatics

1

## Basic Bio-IT Landscape

Overview of our apps and requirements

2

## Hub Stories

War stories from our eval, testing & deployment

3

## <Time Permitting>

Observations, additional details & maybe a promise

4





**QUICK SPRINT: LIFE SCIENCE INFORMATICS**



# Life Science Informatics

- ▶ □ **Biology and Computer Science are a natural fit**
  - □ Atoms, molecules & complex structures such as DNA and Protein easily characterized, stored and represented in ways that computers can read & understand
- ▶ □ **We have 3 core problems:**
  - □ Compute: studying biological data requires sophisticated algorithms, advanced statistical methods and vast amounts of raw computational power
  - □ Storage: Our field is drowning in petabytes of data. Inexpensive lab instruments now routinely *generate terabyte volumes of data per experiment*
  - □ Information Management & Triage: Rate at which we are generating new data exceeds rate at which the storage industry is increasing drive capacity. Something has to give.
- ▶ □ **And one big meta problem**
  - □ More details on that later ...

# The Cliche Example

## “Next Generation” Genome Sequencing

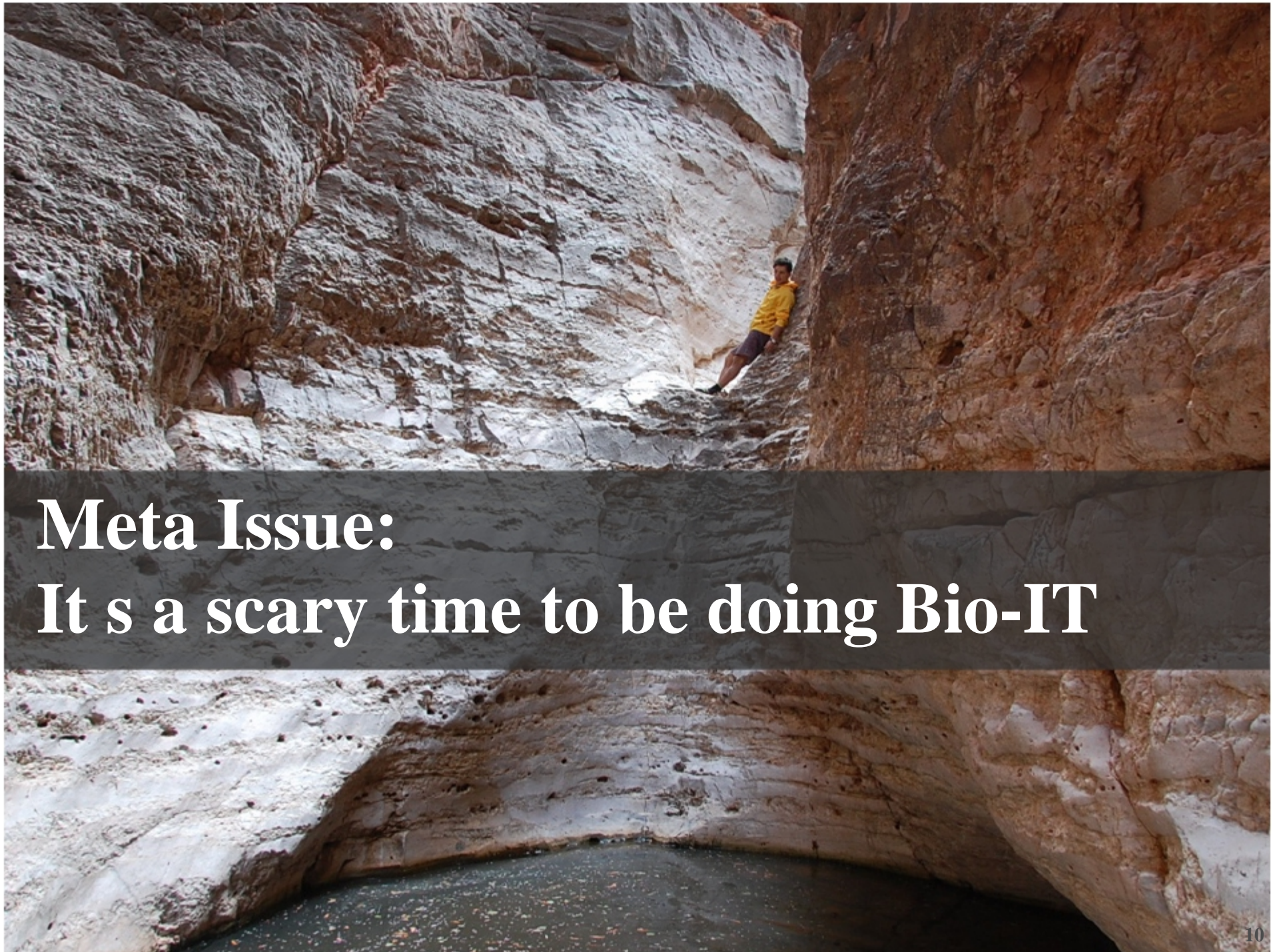
- ▶  **Advances in genome sequencing methods are outpacing Moore's Law**
- ▶  **Since the end of the Human Genome Project, cost-per-base of DNA sequencing has dropped 10,000-fold**
- ▶  **What took an intense global research effort years and billions of \$ can now be done with a \$50,000 instrument sitting on a desktop**



**Illumina MiSeq**

**“Personal Sequencer”**





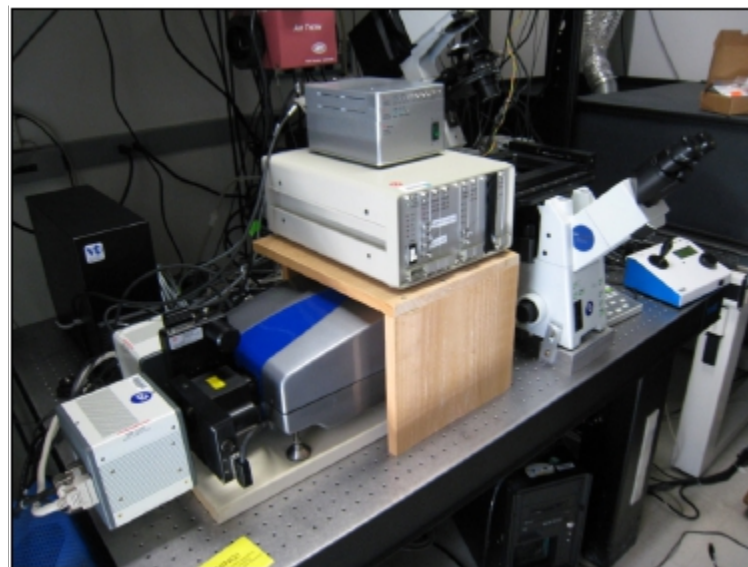
**Meta Issue:**  
**It s a scary time to be doing Bio-IT**



# Big Picture / Meta Issue

This is the driver behind many “Bio-IT” design efforts ...

- ▶ □ **HUGE revolution in the rate at which lab platforms are being redesigned, improved & refreshed**
  - □ Example: CCD sensor upgrade on that confocal microscopy rig just doubled storage requirements
  - □ Example: The 2D ultrasound imager is now a 3D imager
  - □ Example: Illumina HiSeq upgrade just doubled the rate at which you can acquire genomes. Massive downstream increase in storage, compute & data movement needs
  
- ▶ □ **For the above examples, do you think IT was informed in advance?**





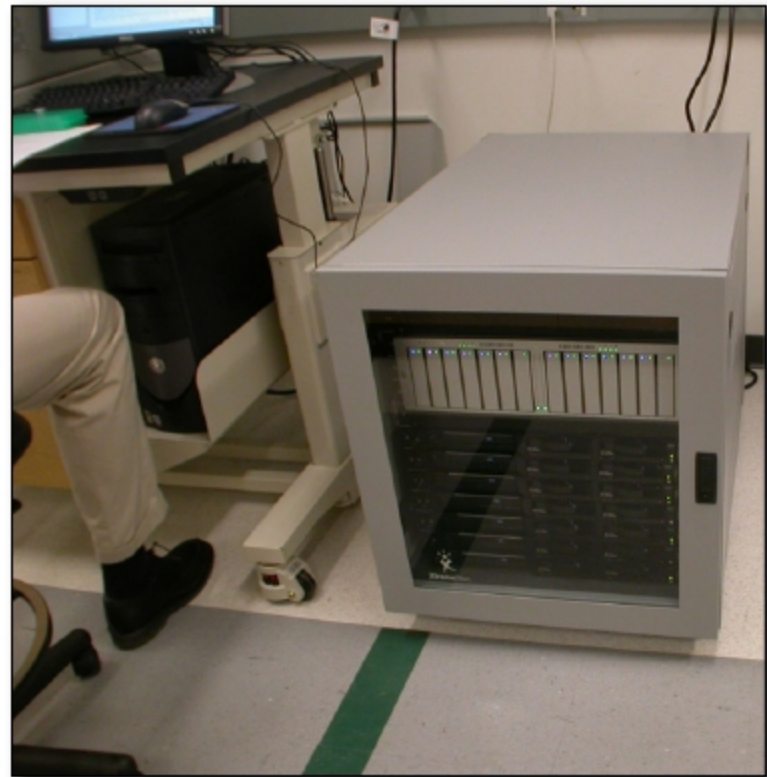
# The Central Problem Is ...

Science progressing way faster than IT can refresh/change

- ▶  **Instrumentation & protocols are changing FAR FASTER than we can refresh our Research-IT & Scientific Computing infrastructure**
  - Bench science is changing month-to-month ...
  - ... while our IT infrastructure only gets refreshed every 2-7 years
- ▶  **We have to design systems TODAY that can support unknown research requirements & workflows over many years (gulp ...)**

# The Central Problem Is ...

- ▶  **The easy period is over**
- ▶  **5 years ago we could toss inexpensive storage and servers at the problem; even in a nearby closet or under a lab bench if necessary**
- ▶  **That does not work any more; real solutions required**







**This is our new normal.**

*Thousands of CPU cores; petabytes of disk*

# And a related problem ...

- ▶ □ It has never been easier to acquire vast amounts of data cheaply and easily
- ▶ □ Growth rate of data creation/ingest exceeds rate at which the storage industry is improving disk capacity
- ▶ □ Not just a storage lifecycle problem. This data *\*moves\** and often needs to be shared among multiple entities and providers
  - □ ... ideally without punching holes in your firewall or consuming all available internet bandwidth

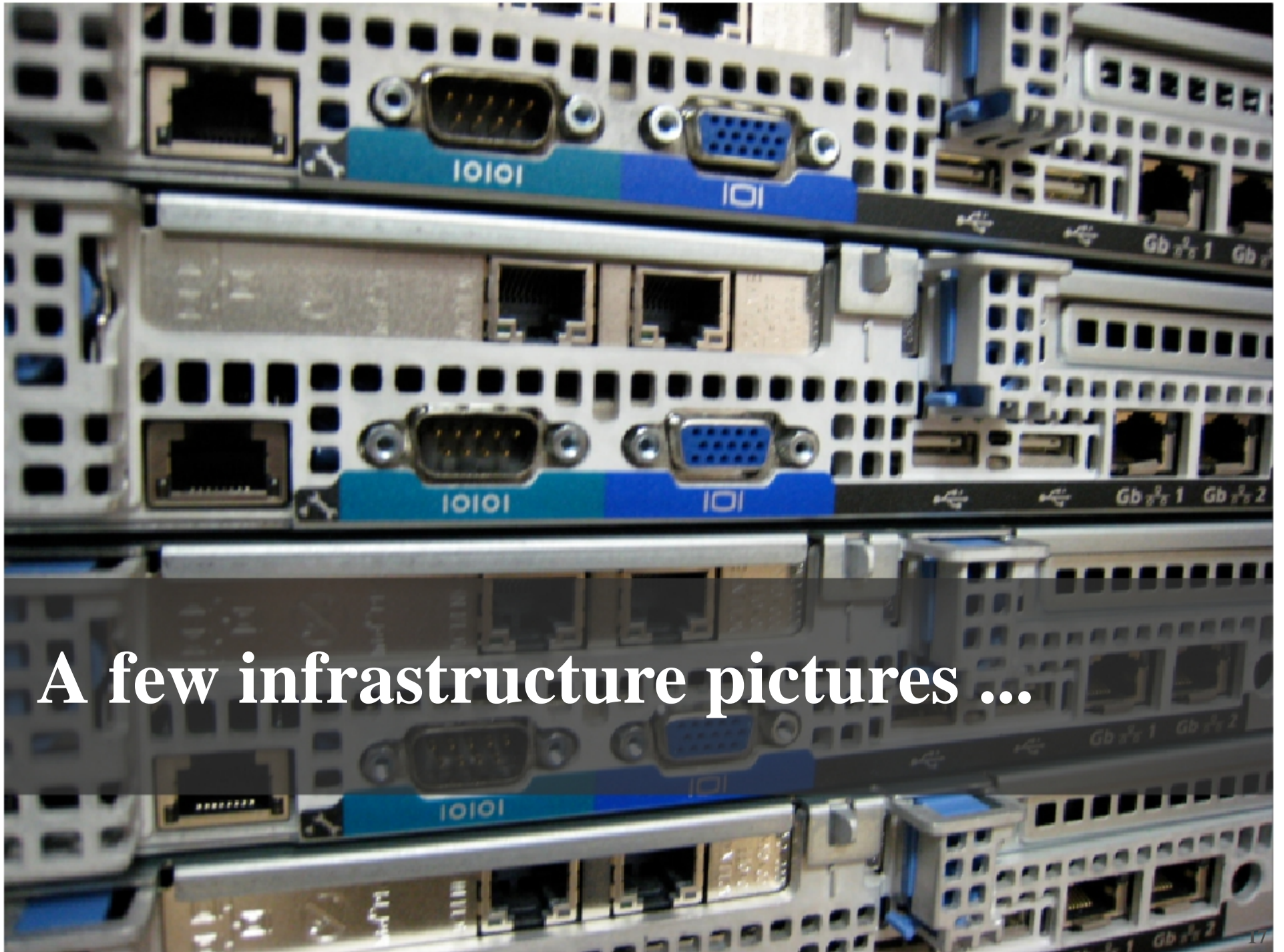




# If we get it wrong ...

The stakes are high.

- ▶  **Lost opportunity**
- ▶  **Missing capability**
- ▶  **Frustrated & very vocal scientific staff**
- ▶  **Problems in recruiting, retention, publication & product development**
- ▶  **And in the clinic: Improper & potentially life-altering medical guidance**



A few infrastructure pictures ...



**The cliché image**





## Lab local HPC & storage

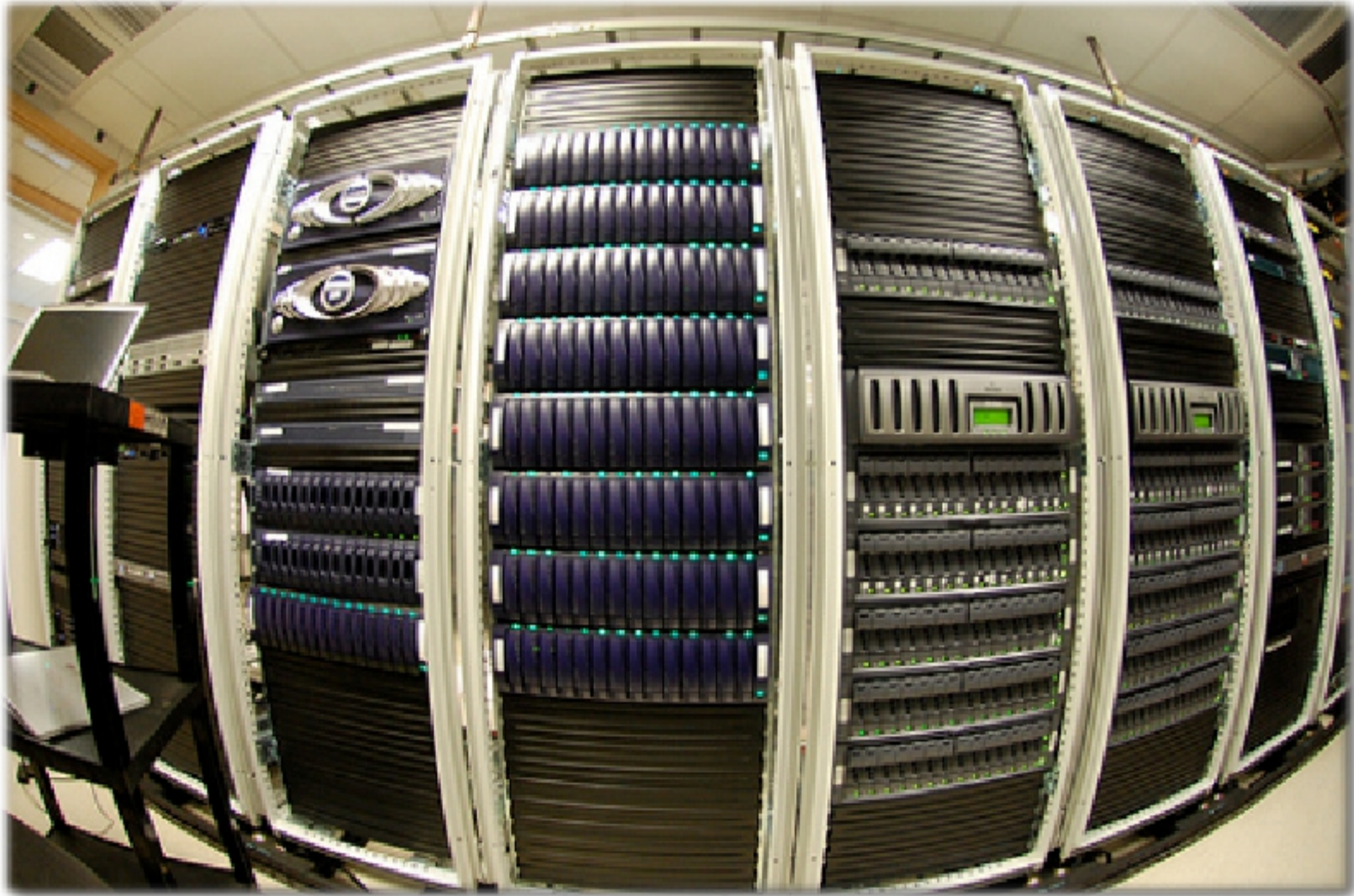


**Small core w/ multiple NGS instrument support**





# Large Core Facility



## **Large Core Facility: Just Storage**





## **Regional Scientific Computing “Hub”**



**Physical data movement station; Unit=  
Naked Disk**





## **Physical Ingest Repo & “Naked” Data Archive**

Name	Instance	AMI ID	Root Dev	Type	State	Status Checks	Alarm S	Monitoring	Security Groups	Key Pair Name	Virtualization
8XL14	i-e622f5ad	ami-e5c9c891	ebs	cc2.8xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
8XL15	i-e422f5af	ami-e5c9c891	ebs	cc2.8xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
8XL16	i-ea29fea1	ami-e5c9c891	ebs	cc2.8xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU1	i-a60cdded	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU2	i-a40cddf1	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU3	i-b80cddf3	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU4	i-be0cddf5	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU5	i-b00cddf6	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU6	i-bc0cddf7	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU7	i-b20cddf9	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU8	i-4a0fde01	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU9	i-b60cddf8	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU10	i-b40cddf	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm
GPU11	i-480fde03	ami-e5c9c891	ebs	cg1.4xlarge	running	2/2 checks passed	none	basic	CST-Studio-EU	EU-nmr-probehead-teamkeys	hvm

**30 of Amazon's largest nodes + 22 GPU nodes: \$30/hr via spot market**

**Yep. This counts.**





# Basic Bio/IT Landscape

# Core Compute

Compute related design patterns largely static

- ▶  **Linux compute clusters are still the baseline compute platform**
- ▶  **Even our lab instruments know how to submit jobs to common HPC cluster schedulers**
- ▶  **Compute is not hard. It s a commodity that is easy to acquire & deploy in 2013**





# File & Data Types

We have them all

- ▶  **Massive text files**
- ▶  **Massive binary files**
- ▶  **Flatfile databases**
- ▶  **Spreadsheets everywhere**
- ▶  **Directories w/ 6 million files**
- ▶  **Large files: 600GB+**
- ▶  **Small files: 30kb or smaller**



# Application characteristics

- ▶ □ **Mostly SMP/threaded apps performance bound by IO and/or RAM**
- ▶ □ **Hundreds of apps, codes & toolkits**
- ▶ □ **1TB - 2TB RAM High Memory nodes becoming essential**
- ▶ □ **Lots of Perl/Python/R**
- ▶ □ **MPI is rare**
  - □ Well written MPI is even rarer
- ▶ □ **Few MPI apps actually benefit from expensive low-latency interconnects\***
  - □ \*Chemistry, modeling and structure work is the exception





# Storage & Data Management

- ▶  **LifeSci core requirement:**
  - Shared, simultaneous read/write access across many instruments, desktops & HPC silos
- ▶  **NAS = easiest option**
- ▶  **Scale Out NAS products are the default standard**
- ▶  **Parallel & Distributed storage for edge cases and large organizations**



# Storage & Data Management

- ▶  **Storage & data mgmt. is the #1 infrastructure headache in life science environments**
- ▶  **Most labs need *peta capable* storage due to unpredictable future**
  - Only a small % will actually hit 1PB
  - Often forced to trade away performance in order to obtain capacity
- ▶  **Object stores and commodity Nexentastor-style methods are making significant inroads**





# Data Movement & Data Sharing

- ▶  **Peta-scale data movement needs**
  - Within an organization
  - To/from collaborators
  - To/from suppliers
  - To/from public data repos
- ▶  **Peta-scale data sharing needs**
  - Collaborators and partners may be all over the world



# We Have Both Ingest Problems

## Physical & Network

- ▶ □ **Significant physical ingest occurring in Life Science**
  - □ Standard media: naked SATA drives shipped via Fedex
- ▶ □ **Cliche example:**
  - □ 30 genomes outsourced means 30 drives will soon be sitting in your mail pile
- ▶ □ **Organizations often use similar methods to freight data between buildings and among geographic sites**





# Physical Ingest Just Plain Nasty

- ▶  **Easy to talk about in theory**
- ▶  **Seems easy to scientists and even IT at first glance**
- ▶  **Really really nasty in practice**
  - Incredibly time consuming
  - Significant operational burden
  - Easy to do badly / lose data



# Huge Need For Network Ingest

1. □ Public data repositories have petabytes of useful data
2. □ Collaborators still need to swap data in serious ways
3. □ Amazon becoming an important repo of public and private sources
4. □ Many vendors now deliver to the cloud



# Cloud(s)

Mainstream in life science for quite some time

- ▶  **IaaS clouds offer excellent pressure release valve when rapidly changing scientific requirements can't be satisfied by on-premise infrastructure**
- ▶  **Economics can't be ignored**
- ▶  **Popular meeting ground for data swapping and collaboration**
- ▶  **Data providers pushing cloud delivery over physical media**
- ▶  **Interesting AWS use cases for archive and downloader pays methods**

```
#!/bin/sh
knife ec2 server create \
-d chef-full \
-N hubzeroTestNode \
-f t1.micro \
--image ami-4d20a724 \
--security-group dag-HZ \
-i ~/bioteam/bioteam-IAM-admins-v1.pem \
--ssh-key bioteam-IAM-admins-v1 \
-x admin
```



# Cloud Hubs & Portals

The 'neutral' meeting ground ..

- ▶  **Many types of entities need to meet, collaborate and exchange life science data**
- ▶  **Data sharing hubs and portals becoming popular on public IaaS clouds like AWS**
- ▶  **Why?**
  - Far easier than punching holes in your firewall and issuing VPN credentials to outsiders



# Cloud Data Repositories

Compelling economics

- ▶  **IaaS clouds becoming center of gravity for large scale scientific data hosting**
- ▶  **Why?**
  - Very compelling pricing
  - You don't pay for the bandwidth used by consumers of your data
  - AWS has some very interesting downloader pays models that seem to be a good fit for grant-funded science with mandated multi-year data accessibility requirements



[www.1000genomes.org](http://www.1000genomes.org)

**It all boils down to this ...**



# Life Science In One Slide:

- ▶  **Huge compute needs but not intractable and generally solved via Linux HPC farms. Most of our workloads are serial/batch in nature**
- ▶  **Ludicrous rate of innovation in lab drives a similar rate of change for our software and tool environment**
- ▶  **With science changing faster than IT, emphasis is on agility and flexibility - we'll trade performance for some measure of future proofing**
- ▶  **Buried in data. Getting worse. Individual scientists can generate petascale data streams.**
- ▶  **We have all of the Information Lifecycle problems: Storing, Curating, Managing, Sharing, Ingesting and Moving**

## Color xterm

```
wgx9@localhost:~$
wxg9@localhost:~$
wxg9@localhost:~$ submit -v cluster-sshOnly dagtest
[DAG-DEBUG submitbatchjob.sh (aspen)] DETECTED NON SGE SCRIPT EXECUTION REQUEST (./00000141_01.sh)
[DAG-DEBUG submitbatchjob.sh (aspen)] SGE will not run jobs that start with digits
[DAG-DEBUG submitbatchjob.sh (aspen)] Will try to rename script to ./sge00000141_01.sh inside /home/hubzero-submit/Jobs/1359613267_00000141_01
[DAG-DEBUG submitbatchjob.sh (aspen)] Will execute command:
  qsh /home/hubzero-submit/Jobs/1359613267_00000141_01/./sge00000141_01.sh
[DAG-DEBUG submitbatchjob.sh (aspen)] Job output below this line :

node22
wxg9@localhost:~$
wxg9@localhost:~$
wxg9@localhost:~$
wxg9@localhost:~$ submit -v cluster-sshOnly sleep60
[DAG-DEBUG submitbatchjob.sh (aspen)] DETECTED NON SGE SCRIPT EXECUTION REQUEST (./00000142_01.sh)
[DAG-DEBUG submitbatchjob.sh (aspen)] SGE will not run jobs that start with digits
[DAG-DEBUG submitbatchjob.sh (aspen)] Will try to rename script to ./sge00000142_01.sh inside /home/hubzero-submit/Jobs/1359613276_00000142_01
[DAG-DEBUG submitbatchjob.sh (aspen)] Will execute command:
  qsh /home/hubzero-submit/Jobs/1359613276_00000142_01/./sge00000142_01.sh
[DAG-DEBUG submitbatchjob.sh (aspen)] Job output below this line :

I appear to be running on node; node22 Current time: Wed Jan 30 20:22:06 EST 2013 I am about to sleep for 60 seconds Current time:Wed Jan 30 20:23:06 EST 2013 \nI am about to sleep for 60 seconds Done. Tool script exiting
wxg9@localhost:~$
wxg9@localhost:~$ █
```

# HUBs for Life Science Informatics



#1

#2

#3

xterm

8:23 Jan 30



2

Firefox [5]

# hubzero and one .gov

## My Hub Story

- ▶  **Sub-contractor to much larger company that won HPC and scientific support contract for division of a very large US .gov disease and public health organization**
- ▶  **Our team has very a specific technical/HPC & scientific computing support mission**
  - ... and a general mission to enhance collaboration, data and knowledge sharing very diverse group of dedicated professionals.
- ▶  **HZ recommended by a team member and supported by senior internal sponsor**



# hubzero and one .gov, cont.

## My Hub Story

- ▶  **I was part of a group that did**
  - Initial evaluation
  - Initial trial deployments
  - Technical assessment report for the go|no-go decision
  - resources required to operate report
  
- ▶  **Along the way ...**
  - Made ugly hacks to Submit() framework to enable integration with Grid Engine managed HPC Clusters
  - ... met a few HZ team members virtually

# hubzero and one .gov, cont.

## My Hub Story

- **And in my free time (non-work) ...**
  - Started work on [Opscode Chef Cookbook](#) for automated installation of Hubzero onto AWS Cloud and other automated DevOps environments
  - <http://bioteam.net/2012/12/hubzero-on-the-cloud/>

```
#!/bin/sh

knife ec2 server create \
  --verbose \
  --template-file /Users/dag/opscode-platform-projects/bioteam/chef-repo/.chef/bootstrap/debian6-gems.erb \
  --availability-zone us-east-1b \
  --security-group dag-HZ \
  --node-name hubzeroTestNode \
  --flavor t1.micro \
  --image ami-4d20a724 \
  --ssh-user admin \
  --identity-file /Users/dag/opscode-platform-projects/bioteam/bioteam-IAM-admins-v1.pem \
  --ssh-key bioteam-IAM-admins-v1
```

# Current Status

Hubzero and .gov story

- ▶  **In production & formally deployed**
  - ... but in small-scale pilot mode
- ▶  **Successful integration with Grid Engine and our HPC cluster & storage environment**
- ▶  **Primary current use is documentation**
- ▶  **Secondary focus will be Collaboration/Projects**
- ▶  **Beginning to recruit users and encourage the more social collaboration and self-publishing features**



# Future Status

Our sweet spots and interest areas ...

- ▶  **Very significant interest in courseware & learning**
- ▶  **Very significant interest in the DataStore features**
- ▶  **Interest in tags, especially automated tagging**
- ▶  **Interest in the fine-grained security model & role-based access to content**
- ▶  **Interest in automated/API-driven content creation. Possibly by lab instruments that auto publish experimental result summaries w/ links to data repository location**



Time permitting ...

*image: [shanelin](#) via flickr*

# Time?

A few more details

1.  My take on our Hubzero assessment
2.  Enterprise integration war stories
3.  Potential Personal Promise



# My \$.02: Internal Hubzero assessment

Speaking for myself, not my employer or .gov team !

## ▸ □ **Initial Impressions**

- □ Very positive
- □ The public hubs are doing a great job at evangelizing
- □ Particularly good job at delivering a CMS that understands the org structure and working behaviors of real scientists
- □ Great foundation for what we were looking to do

## ▸ □ **Observed, however:**

- □ Not many commercial/industry users publicly visible
- □ HZ not easily deployed in enterprise environments (OS/vm/etc)
- □ The most sophisticated Hubs appear to be hosted at Purdue where they depend on internal expert support and may also leverage knowledge that otherwise may not be well documented

# My \$.02: Internal Hubzero assessment

Speaking for myself, not my employer or .gov team !

- ▶  **Assessment summary outcome:**

- Worth pursuing; great features and capabilities
- IT Infrastructure requirements for running a hub are trivial relative to the stuff we have already supporting peta-scale genomics

- ▶  **However:**

- Human resource requirements were non trivial and if senior management was serious about deploying a hub in any meaningful way they would have to commit real dollars and real staff to the effort.

# My \$.02: Internal Hubzero assessment

Speaking for myself, not my employer or .gov team !

- ▶  **Human resources required for a successful Hubzero deployment:**
  - Salaried FTE or part-time internal employee (NOT AN OUTSIDE CONTRACTOR) required to build community, configure the Hub and serve as Community Manager
  - Need a Web Developer w/ creative skills and UI experience (the best hubs have great UI and visuals)
  - Need a Joomla CMS expert on contract or on-call
  - Existing IT staff would need C++/Rapture training

# War Stories

Episodes that drove the assessment results

1. □ Outsider vs Insider issues
2. □ Enterprise deployment issues
3. □ Authentication & identity management
4. □ Featured Article adventures
5. □ Submit() and Rapture adventures



**One last thing ...**

# One last thing

- ▶  **In 2013 when I can ...**
  - ... Launch a completely integrated 500 CPU/GPU cluster on AWS with 1 command typed from my macbook
  - ... Launch a full Wordpress stack with global CDN, SSL-aware load balancers, geographically replicated SQL datastores and an elastic fleet of web servers via a single CloudFormation .json template
- ▶  **It s hard to believe that Hubzero is not available for everyone as a single button click on Amazon or Google Compute**

# One last thing

- ▶  I've benefited personally and professionally from open source over the years; have always tried to contribute back in kind
- ▶  If there is interest in making Hubzero available in forms other than VM files and package-based installs I might be able to contribute
- ▶  **Personal interests include:**
  - Packaging Hubzero for [AWS Marketplace](#) (\$0.00 product)
  - Creating [Opscode Chef](#) Cookbook that can auto build and deploy Hubzero from packages on any Debian/Ubuntu OS

