# Maintaining a publicly used analysis infrastructure



Keeping everything running when everyone's an analyst

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#### Where we were, where we're going

In the past	In the future
Analysts did analysis	Many people do analysis
A small group of people shared computational resources	More people sharing, more diverse needs
Communication about and tracking load on servers/clusters is fairly straightforward	more complex – more groups, more people involved
Users pretty comfortable with Linux	Training and mentoring is needed

## **BRACE YOURSELVES**

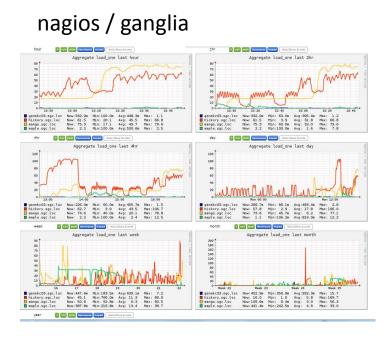


## My institute's (current) approach to sharing computational resources

- Most servers / clusters are currently open
- A few are dedicated to particular groups
- Some clusters require training to get access
- We made "server usage guidelines"
  - Use < half of available memory and cores on a server</li>
- Encourage use of htop
- Informal "Policing" via email

### Challenges we've had

- Encouraging people to be mindful, use htop
- Resource hog programs, novice users
- When do we add computational resources?



#### Analyst grumbling

#### March 2014 Archives by thread

- Messages sorted by: [subject] [author] [date]
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Starting: Mon Mar 3 11:02:08 CST 2014 Ending: Fri Mar 28 15:46:36 CDT 2014

- [linux-discuss] hickory Paulson, Ariel
  - [linux-discuss] hickory McGee, Jenny
    - [linux-discuss] hickory McGee, Jenny
    - [linux-discuss] hickory Meier, Samuel [linux-discuss] hickory McGee, Jenny
- · [linux-discuss] Hickory Paulson, Ariel
- o [linux-discuss] Hickory Dietz, Dustin
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  - [linux-discuss] Hickory Dietz, Dustin
- [linux-discuss] Color Exhibit Gogol, Madelaine [Linux-power-users] desktop reboot Miller, Danny
- o [Linux-power-users] desktop reboot Seidel, Christopher
- o [Linux-power-users] desktop reboot Paulson, Ariel
- [linux-discuss] circos installation | Seidel, Christopher [linux-discuss] circos installation Gogol, Madelaine
  - [limux\_discuss] circos installation | Spidal Christopha

#### Other Analysis Resources

- Genomes, annotations, alignment indexes
- Databases (e.g. GO, BLAST)
- Software Installation, Updates
- Public Sequencing Data
- Galaxy
- Rstudio Server

#### How about you?

- Balancing user access vs. user responsibility
- Breakdown of "private" versus "public" resources
- Challenges faced and solutions
- Tracking load, deciding when to add resources
- Managing genomes/software/data

