ABCD-LMA, August 16, 2018

"Why Do Medical Students Need a 10,000-Core Cluster?"

Questions about HMS Research Computing

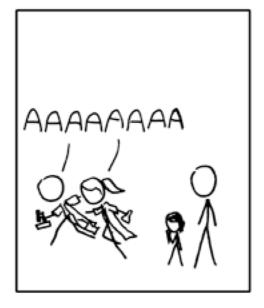
Amir Karger

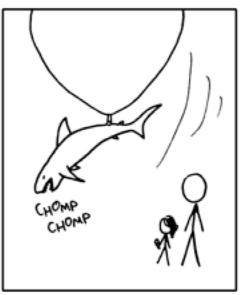
Research Computing Group
Harvard Medical School Information Technology

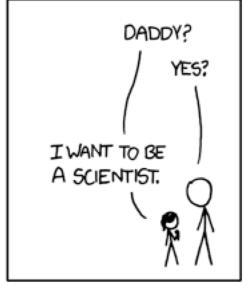
Whom Does Research Computing Support?

- 700 MD students
 - Few do research, so who needs RC?
- 800 PhD students doing basic research
- 1500 post-docs "on Quad"
- Thousands more:
 - HSDM, HSPH
 - 16 Affiliate hospitals
 - Collaborators around the world

What Does RC Support?







XKCD #585

Science!

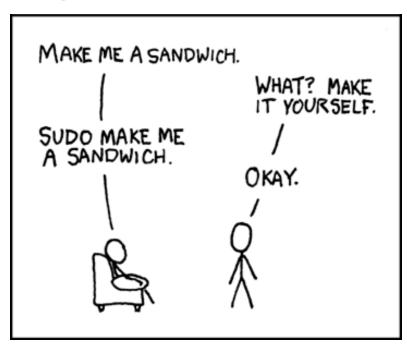
What Does RC Support II?

- Basic research: bacteria, worms, mice
- Football Players' Health Study (HIPAA!)
- Tracing drug-resistance in tuberculosis
 - Farhat Lab, GHSM + DBMI
- Opiods: duration more serious than dose
 - Kohane Lab, Biomedical Informatics
- Mapping neurons flies use to smell
 - Lee Lab, Neurobiology
- Pretty pictures at the end

Who Are You?

DevOps (5 people)

- HPC cluster administration
- Automating sysadmin ("big red lever")
- Build Level 4 systems
- OMERO
- SQL and web hosting



Who Are You II?

RCCs – Research Computing Consultants (6)

- Science degrees
- Programming experience
- Knowledge of HMS computing environment

What Are You Doing?!

- Consulting! Listen, think, advise
- Science better, cheaper, faster
 - Use the right resources, programs, statistics
 - Run workflows in the most efficient way
 - Scale up workflows
- Most services are free (!)
 - We (almost) never analyze data for users
 - We show users how to do it
 - Or send to other groups (like HCBC)

How Can We Empower Researchers?

- Desktop/workstation -> O2
- Single gene analysis -> thousands
- Parallelization, GPU
- R, Perl, Python, bash, programming
- SQL Database hosting
- Hosting research websites
- Data manipulation hacking wrangling science
- Do bioinformatics, biostatistics correctly

How Can RC Scale?

- Showing is more efficient than doing
- Hands-on classes lasting about 2 hours
 - Intro to Linux & O2
 - Introduction to NGS Analysis
 - Perl, Python
 - R/Biostatistics and Bioconductor
 - Parallel Computing
 - Matlab
 - git and Github
 - Ad-hoc classes, bigger rooms as needed

What is Your Quest?

Orchestra: entered production 2004

- 5000 cores 7+ years old, 12-core nodes, 96 GB RAM
 - (+2000 faster cores, 256 GB RAM)
- Old LSF scheduler
 - \$2M upgrade + ongoing support needed
 - No help in Google
 - Lacking features priority bumps difficult
- 12 years of clutter
 - 250 queues, old software versions, config...

What is Your Quest II?

O2 went into production 9/2017

- 5000 new cores (+ 2000 stolen from Orchestra)
 - Fast-ish processors, 32-core nodes, 256 GB RAM
- Slurm scheduler:
 - Open, popular
 - Free, with cheap support
 - Feature-ful more knobs and dials
- 12 partitions
- New OS/software, puppet, ...

What's the Difference?

New scheduler commands

- bsub -q short -W 1:0 myprog in.txt ->
 sbatch -p short -t 1:0:0 myprog.batch in.txt
- bjobs -u ak150 -> squeue -u ak150
- New paths for some filesystems
- New module names & versions
- Some new libraries
- Users need to change commands, scripts, ...
 - How hard could it be to convince them?

What's the Worst that Could Happen?

- RC too distracted to build the new system
 - Maintaining 2 systems and migrating users
- Labs refusing to leave Orchestra
 - Can't migrate crucial workflow
- Labs delaying leaving Orchestra
 - "Orchestra works just fine"
- Labs insisting we bring Orchestra back
 - Didn't know about move, didn't test

What Do You Expect?

- Complaining or oblivious users
 - Carrots only encourage 75% of users
 - When do we employ sticks?
- Details missed by busy RC staff
- Timeline:
 - Expectation: 1 year
 - Fear: 18 months? More?
 - Reality: ???

How Do You Get There From Here?

- Minimize differences to lower barriers
 - Same partition names, limits, module "API"
- Notify users early and often
- Document changes fully and concisely
- Run migration workshops
- Hire term employee focusing on migration
- Use every ticket as an opportunity (to pester)
- Move fast nodes to O2

Are We There Yet?

- "All carrots, (almost) no sticks" ... worked!
- ~1000 users migrated in 7 months
 - Not including reeeeeally long beta
- Labs refusing to leave Orchestra: 0
- Orchestra turned off 4/2018 (5000 days)
- Labs asking to bring Orchestra back: 0
- Um, some websites kept running on Orchestra
 - Minimal usage, many nodes turned off

Did You Get Very Far?

- O2 is our flagship service
 - Many types of life science research
 - Batch and interactive work (even GUI!)
- Lots of usage
 - ~7000 faster cores, 32 cores per node, 256 GB
 - GPU & high memory machines
 - 200 users, 70k jobs per day (30% > Orchestra)
 - 3000 cores, more GPU+highmem coming soon
- Efficient usage is still a struggle

What Supports RC?

Storage

- 1 PB Scratch disk on O2
- 1 PB "no_backup" (growing)
- 21/27 PB Tier 1
- 1.5/8 PB Tier 2 (growing a lot)
 - Cloud storage with local "stubs"
- Tier 3 to come
 - Glacier et al.

What Supports RC II?

Network

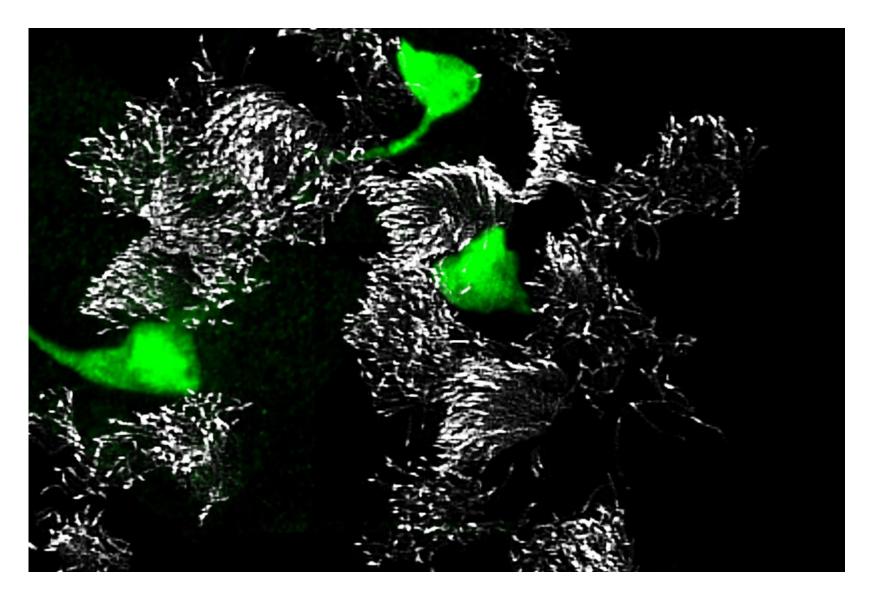
- 1 Gb for end user devices
 - 10 Gb for some microscopes, etc.
- 4 x 10 Gb to building switch, 2 x 100 to core
- Multiple 100 Gb to/between data centers (Markley, Lowell)
- 2 x 10 Gb (only!) to LMAnet -> Intertubes
- 30 TB/day to/from LMAnet, 100 MDC

Who Supports RC?

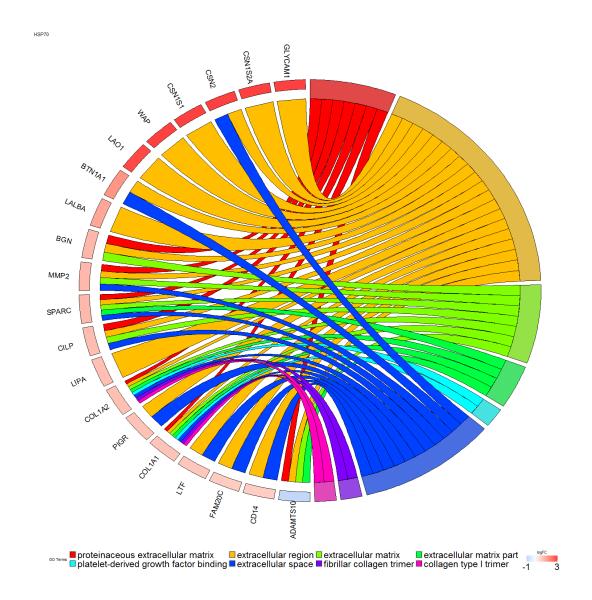
- Infrastructure Ops (IOps) network/storage
- Client Services Group desktop support
- Chan Bioinformatics Core analysis, training
- RITS, SBGrid, WARP, etc. specialized support

The Dean – "hard money" from IT

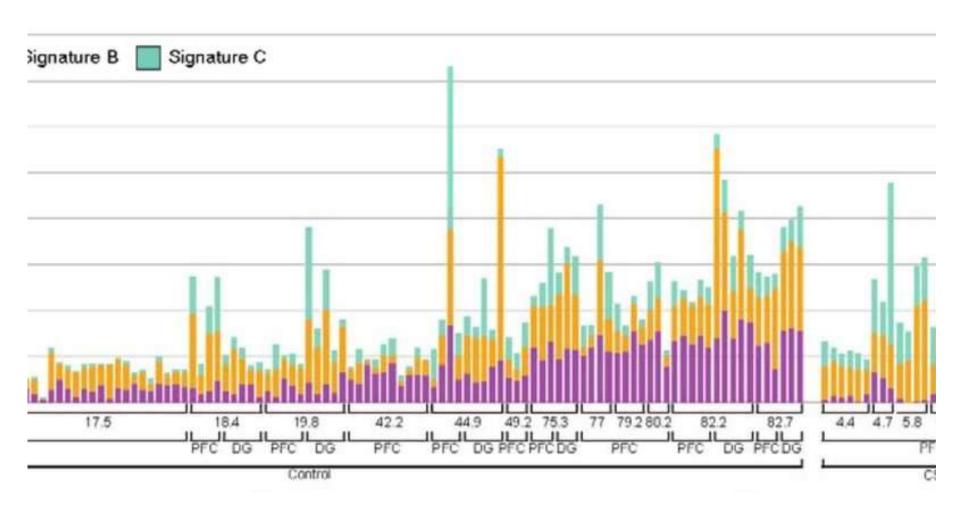
And now ... pretty picture time!



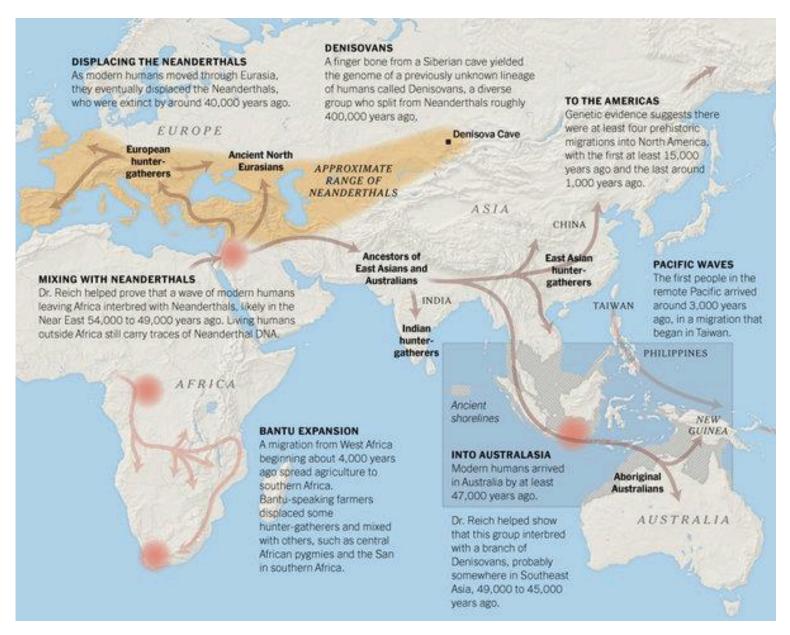
Klein Lab, Systems Biology (Nature, 8/2018)



Calderwood Lab, BIDMC (In press)



Walsh Lab, Childrens Hospital (Science, 12/2017)



Reich Lab, Genetics (New York Times, 3/2018)

Any (of Your) Questions?

http://rc.hms.harvard.edu