

Getting the most from the farm at the Sanger Institute

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Table of contents

Overview of the Sanger

Monitoring

Weekly Farmers Standup Meeting

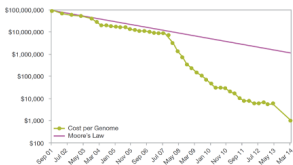
Lustre Reporting

The Campus



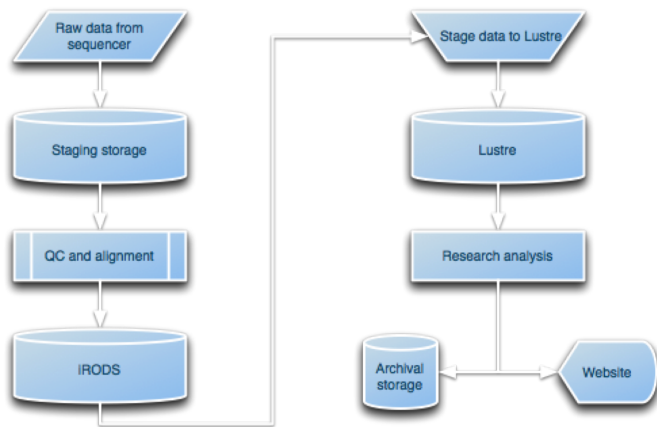
- ▶ About 900 staff at the Sanger
- ▶ 500 at the EBI
- ▶ New sequencing building under construction
- ▶ Expect about 30 spinout and startups on site in 2 years

History



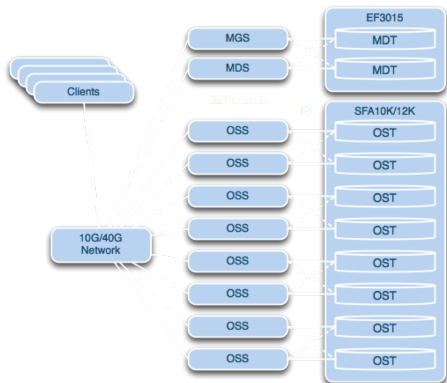
- ▶ Set up in 1993
- ▶ 1998 - Nematode worm Completed (97Mbp)
- ▶ 2003 - Human Genome Complete (2000Mbp)
- ▶ 2004 - MRSA Genome
- ▶ 2005 - Current Data Centre opens
- ▶ 2008 - Next Generation Sequencing, 1000 Genomes Project begins
- ▶ 2009 - Joins International Cancer Genome Consortium
- ▶ 2010 - UK10K Project begins
- ▶ 2013 - UK10K Completed

Typical Workflow

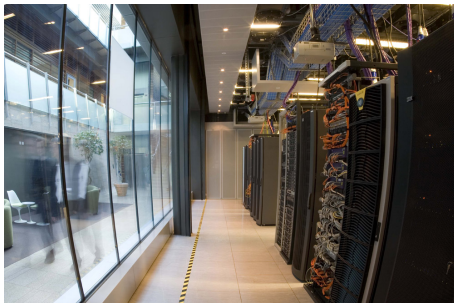


The Cluster

- ▶ 11 Lustre Volumes, 2 more imminent, one to be retired
- ▶ 250TB/500TB/1PB each
- ▶ 6PB total capacity
- ▶ DDN Exascaler hardware
- ▶ Our own lustre software install
- ▶ Aim to deliver 5MB/s for each core
- ▶ IB Connected OSS - OST
- ▶ 10GigE to clients
- ▶ 28PB storage overall (lustre, iRODS, NFS)
- ▶ 17,000 cores of compute - mostly HP Blades



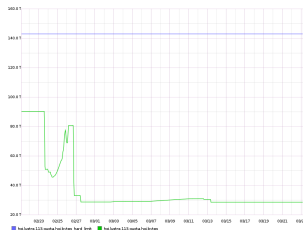
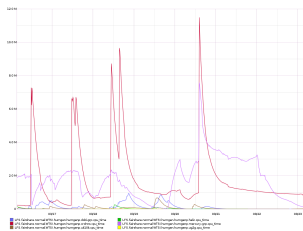
Monitoring



- ▶ Ganglia, Opsview / Nagios, Platform LSF / Tableau Analytics in production
- ▶ Graphite in test / development
- ▶ Hardware ordered for production graphite cluster
- ▶ May switch to InfluxDB
- ▶ Currently collecting standard metrics (cpu, ram, disk, network)

Application Level Monitoring















- ▶ Graphite makes it easy to add metrics
- ▶ Cron which collects Platform LSF Fairshare snapshot every 10 mins
- ▶ Has been useful for identifying cpu accounting kernel bug
- ▶ Also capturing lustre quota information for groups
- ▶ Working on real time analysis for captured data
- ▶ Pre-emptive warn of over-quota



The Weekly Report

- ▶ Generate weekly usage report for Human Genetics
- ▶ python script gets data from LSF Analytics DB (Vertica)
- ▶ Gets jpeg image and user name from ldap
- ▶ Merges the data with a \LaTeX template using pyratemp
- ▶ Generates nicely formatted report
- ▶ A few copies handed out at the meeting

CPU Used since 2015-03-12 : Top 20 Users

	Name	Cores / Weeks	Avg. CPU Efficiency	# Jobs	Avg. Run Time (Hrs.)
	Thomas Costanzo (tdc)	523.03 373.24	6.24 ± 0.73 6.24 ± 2.04	32469 4344	2.71 14.46
	anonymous (anonymous)	1.80 162.91	4.89 ± 0.79 3.75 ± 1.52	67647 148090	0.01 0.31
	Sebastian Singh (ss)	96.75 16.63	5.98 ± 0.58 5.79 ± 0.85	77962 19906	0.22 0.15
	Stephen Scalet (sca)	94.08 8.44	65.04 ± 21.75 133.84 ± 61.21	595 21	2.56 3.15
	Amelia Anton (aant)	18.63 8.65	0.33 ± 2.72 0.24 ± 2.12	17523 10567	3.42 3.60
	Yuan Chen (yuan)	23.25 0.41	6.24 ± 0.19 6.10 ± 0.44	1373 531	2.85 0.13
	Yang Liu (yli)	13.45 6.86	6.50 ± 7.37 6.19 ± 4.59	144 62	15.07 18.77
	Overlapping Developmental Disorders (ODD)	7.65 12.38	2.48 ± 2.52 5.99 ± 1.88	2277 671	1.42 3.23
	Christopher Franklin (cfr)	13.40 0.01	6.12 ± 0.43 6.18 ± 1.48	46 24	50.00 0.08
	Andrei Bogdanov (abd)	12.96 0.10	11.12 ± 5.34 3.35 ± 5.84	486 105	2.52 0.29
	Stephan Schiffels (sch)	8.96 3.92	22.99 ± 9.43 19.70 ± 9.35	307 190	1.33 1.10
	Nick Williams (nwi)	8.77 0.00	31.46 ± 13.79 0.00 ± 0.00	66 0	4.44 0.00
	Michael Spink (msp)	7.86 0.00	6.11 ± 0.10 0.00 ± 0.00	13 0	104.03 0.00
	Spice Monk (sm)	0.12 4.72	6.28 ± 0.06 6.18 ± 2.70	2 7	9.65 114.56

The Meeting

- ▶ Human Genetics get together over coffee every thursday at 3.
- ▶ Send out the report and try to get at least the top 10 to turn up.
- ▶ Stand in a circle and each user says...
 - ▶ What they were doing - the science
 - ▶ Job submission strategy
 - ▶ queue
 - ▶ how many jobs
 - ▶ memory requirements
 - ▶ threading
 - ▶ Any problems ?
- ▶ Trying to get an idea of what the best practice is
- ▶ Identifying areas where
 - ▶ We need more documentation
 - ▶ May need to improve the systems - any kernel / lustre bugs etc.
 - ▶ We could improve tools available to users
- ▶ Great for building a community

The Constraints

- ▶ Team / project users share quota
- ▶ Users can be members of multiple projects
- ▶ Project lifetime longer than user tenure
- ▶ Long-term need for intermediate project data
- ▶ Due to proliferation of projects, quotas overprovisioned
- ▶ We need users to tidy up after themselves
- ▶ Users need to know where the data is

The Problem

- ▶ I need to run some analyses, how much space is available for use by my project?
- ▶ You asked us to clean up the disk, where are the oldest large files so I can prioritize them for archiving or deletion?
- ▶ My project is near the quota limit. Where is all the space being used ? Who is using it ?
- ▶ This is not easy...
 - ▶ lfs quota - gives usage but no idea where files are
 - ▶ lfs find - stops as soon as you hit "permission denied"
 - ▶ find - very hard on the MDS, syntax tricky for users
 - ▶ df - can see usage but no granularity
 - ▶ du - continues through "permission denied", hits MDS hard, slow, difficult for users
 - ▶ agedu - data collection takes a very long time, updated rarely, large list of files in order of last accessed

Towards a solution - mpistat

- ▶ Guy Coates found paper on efficient parallel file tree walking using MPI
- ▶ Implemented the algorithm with a python class
- ▶ Subclassed the walker to make a fast parallel copy program
- ▶ I made it do an lstat instead.
- ▶ Get full lstat for an entire volume in a practical amount of time (tens of minutes).
- ▶ Difficulty in formatting the output - file names with unprintable characters
- ▶ Solved by base 64 encoding the path in the tab formatted output file

Summary Report

- ▶ Ballpark estimate of cost to store file - £150 per Terabyte per Year
- ▶ Calculate a cost for every file based on size and a time
 - ▶ ctime - cost to store since creation
 - ▶ atime - cost to store since last access - i.e. wastage
- ▶ Keep tally of following - totals, by user and by group
 - ▶ file sizes
 - ▶ file counts
 - ▶ zero length files
 - ▶ inode type - how many files / directories / symlinks
 - ▶ costs
 - ▶ files with unprintable characters
 - ▶ Example report...

Lustre Treemaps

- ▶ Summary all well and good
- ▶ Want users to be able to interact with the data
- ▶ Treemaps highly suitable
- ▶ C++ program...
 - ▶ Parses the mpistat output
 - ▶ Builds in memory tree, node for each directory
 - ▶ Keeps track of accumulators for summaries at each node
 - ▶ Embedded http server using facebook proxygen framework
 - ▶ Can GET json representation of the tree
- ▶ Web Frontend
 - ▶ Queries the tree for json of particular subtrees to a given depth (usually 3)
 - ▶ Renders treemap using d3.js
- ▶ Demo...

Performance and Future Plans

Volume	Files	Size	mpistat	tree build	RAM
scratch114	5.5M	769TiB	23m	17m	5GB
scratch111	16.6M	276TiB	10m	54m	20GB
scratch113	34.6M	649TiB	69m	100m	61GB

- ▶ Need gzip encoding of response
- ▶ Speed up treebuild - multithreading
- ▶ Use key-value store instead of RAM - lmbd
- ▶ Real-Time updates - tap into lustre changelog mechanism
- ▶ Or use Robin-Hood

Acknowledgements and References

- ▶ Peter Clapham - Platform LSF / Tableau Analytics / Vertica
- ▶ Simon Fraser - Local Graphite Guru
- ▶ James beal - Resident Lustre Expert
- ▶ Matthew Rahtz - Grafana wiz, git-foo
- ▶ John Constable - Systems team presence at farmers standup
- ▶ Tim Cutts - Sanger overview slides
- ▶ Guy Coates - parallel filetree walker, parallel copy
- ▶ Parallel Filetree Paper
- ▶ Parallel Filetree Website
- ▶ Josh Randall - lustre tree front end
- ▶ Martin Pollard - investigating lustre changelog mechanism
- ▶ HGI Github