

Computational infrastructure for NGS data analysis

Pablo Escobar
pescobar@cipf.es

Computational infrastructure for NGS

- In NGS we have to process really big amounts of data, which is not trivial in computing terms.
- Big NGS projects require supercomputing infrastructures

Data tsunami is real

- Some disks in our lab.....



Sequencing cost vs IT cost

Sequencing cost goes down....so IT cost goes up

Full Genome Sequencing & The Genetic Revolution

Cost per Human Genome vs Total Number of Genomes Sequenced



www.existencegenetics.com

Industry data from public online sources

■ Cost per Human Genome for Full Genome Sequencing

■ Total Number of Human Genomes Sequenced

Dashed lines represent extrapolations based upon current trends

Computational infrastructure for NGS

These infrastructures are expensive and not trivial to use, we require:

- Conditioned data center (servers room). This is expensive
- Computing cluster:
 - Many computing nodes (servers)
 - High performance and high capacity storage
 - Fast networks (10Gb ethernet, infiniband...)
- Skilled people in computing (sysadmins and developers).
 - In CNAG currently 30 staff - >50% informatics

Computing cluster

- Distributed memory cluster
 - 8 or 12 cores by node
 - x86_64 arch
 - At least 48GB ram per node
- Fast networks
 - 10Gbit
 - Infiniband
- Batch queue system (sge, condor, pbs, slurm)
- Many GPUs tools are being developed, no a bad idea to have some if you plan to use gpu tools



Storage system

- Storage is the most important piece in the IT infrastructure for NGS
- Storage is the most expensive
- Good design is really important. Talk with experts
- Keep in mind the storage scalability.
- Try to keep storage flexible. Changes come fast

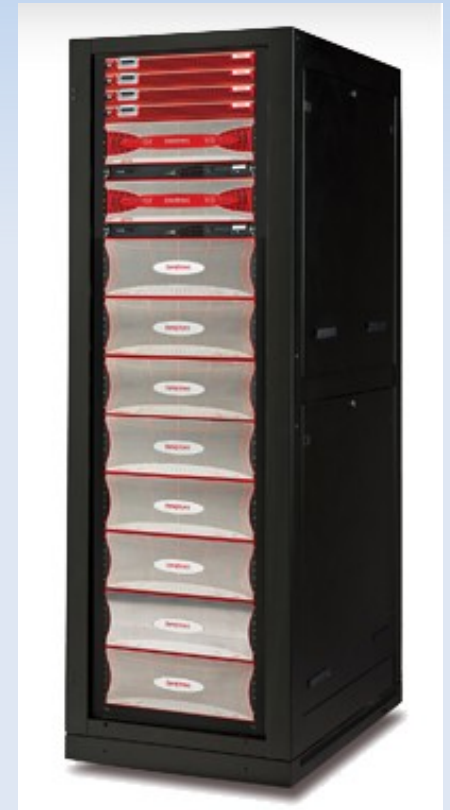
Storage System

- Traditional backups are a problem, if even possible.
- Raid is your friend.
- Plan a good data storage policy
- Recommended reading:

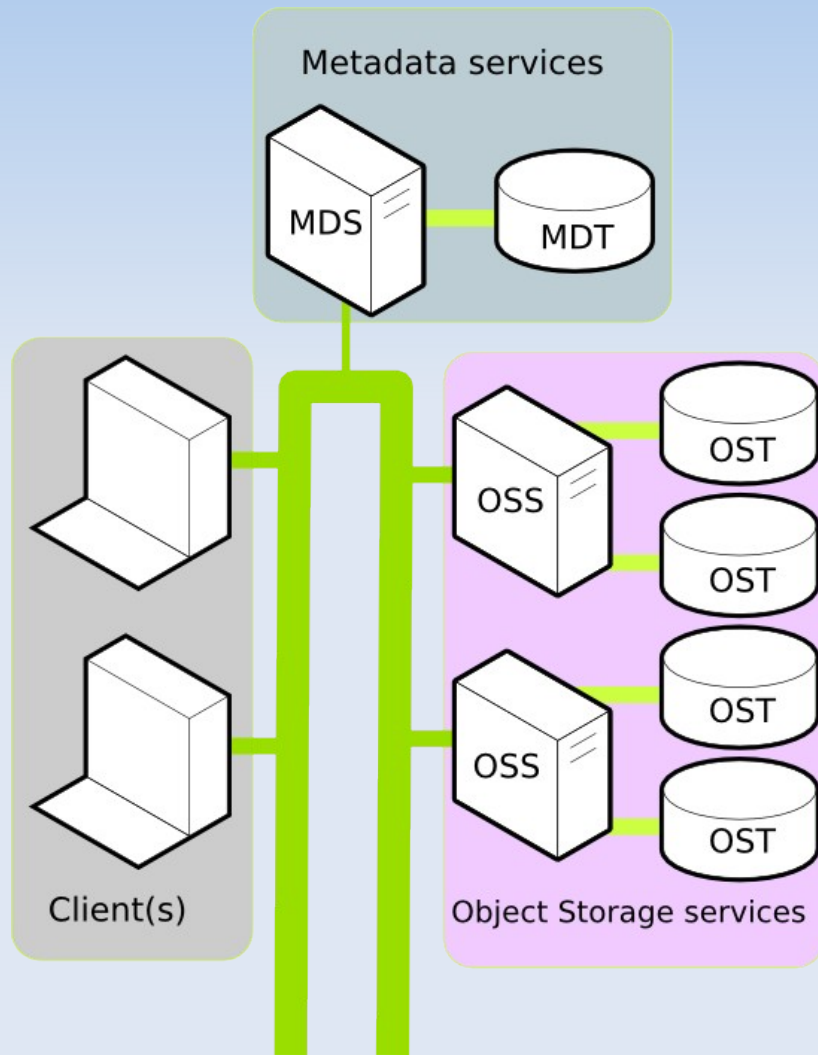
http://www.bioteam.net/wp-content/uploads/2010/03/cdag-xgen-storageForNGS_v3.pdf

Storage system

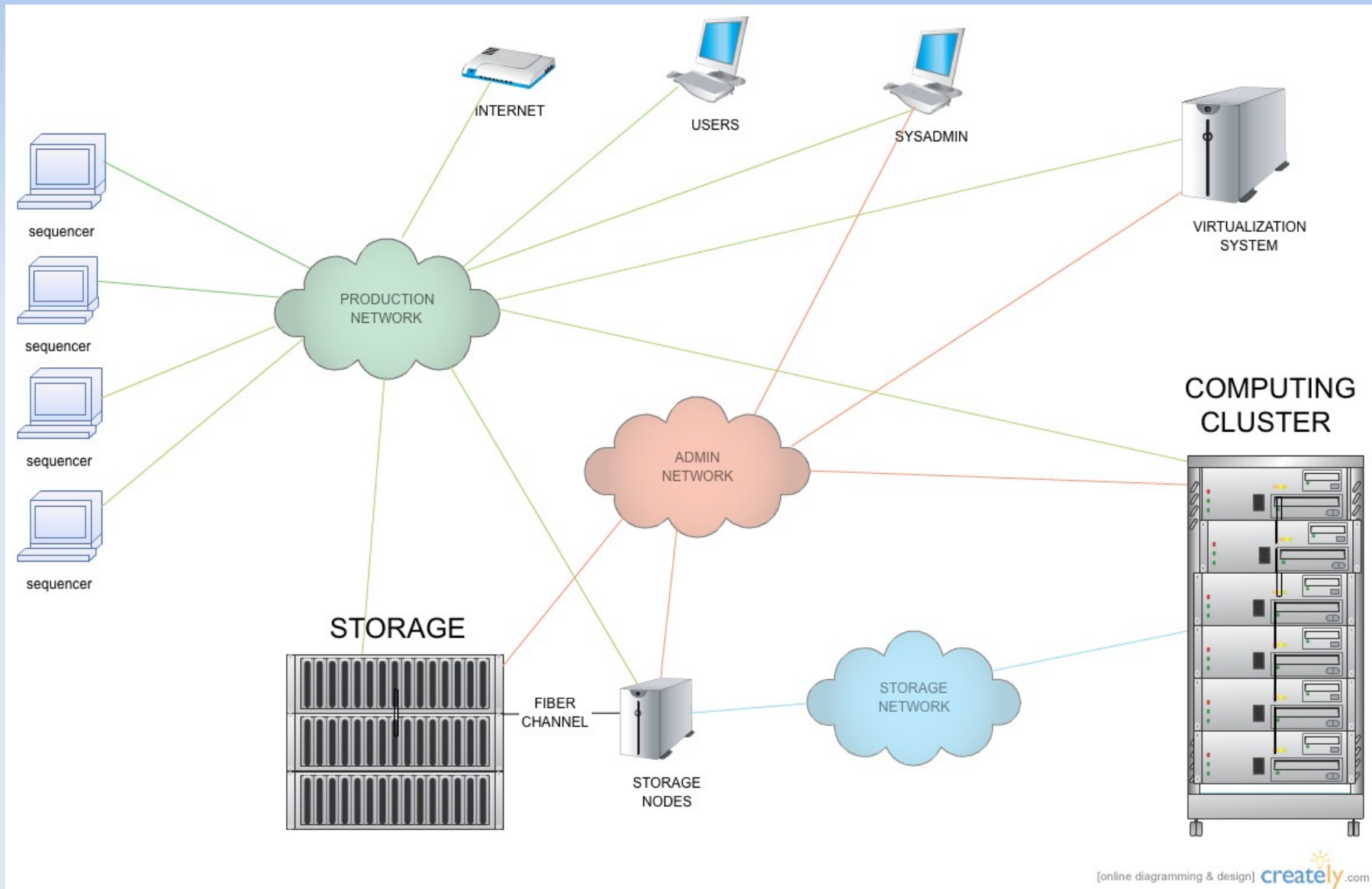
- Distributed filesystem for high performance storage
 - Lustre
 - GPFS
 - Ibrix
 - GlusterFS
 - Panasas
 - Isilon
- These filesystems are not trivial to administer
- NFS is not a good option for supercomputing



Distributed filesystem schema



Infrastructure schema



Small infrastructure

- Recommended at least 2 machines
 - 8 or 12 cores each machine.
 - 48Gb ram minimum each machine.
 - BIG local disk. At least 4TB each machine
 - As much local disks as we can afford
- Price range: starting at 8.000€ - 10.000€ (two machines)

Sequencing centers in Spain

Medical Genome Project

- Sequencing Instruments
 - 7 GS-FLX (Roche)
 - 4 Solid™ 5500 (Applied Biosystems)
- Informatics infrastructure
 - 300 core cluster
 - 0,5 petabyte ibrix filesystem

Medical genome project

Storage racks



IBRIX filesystem
front-ends

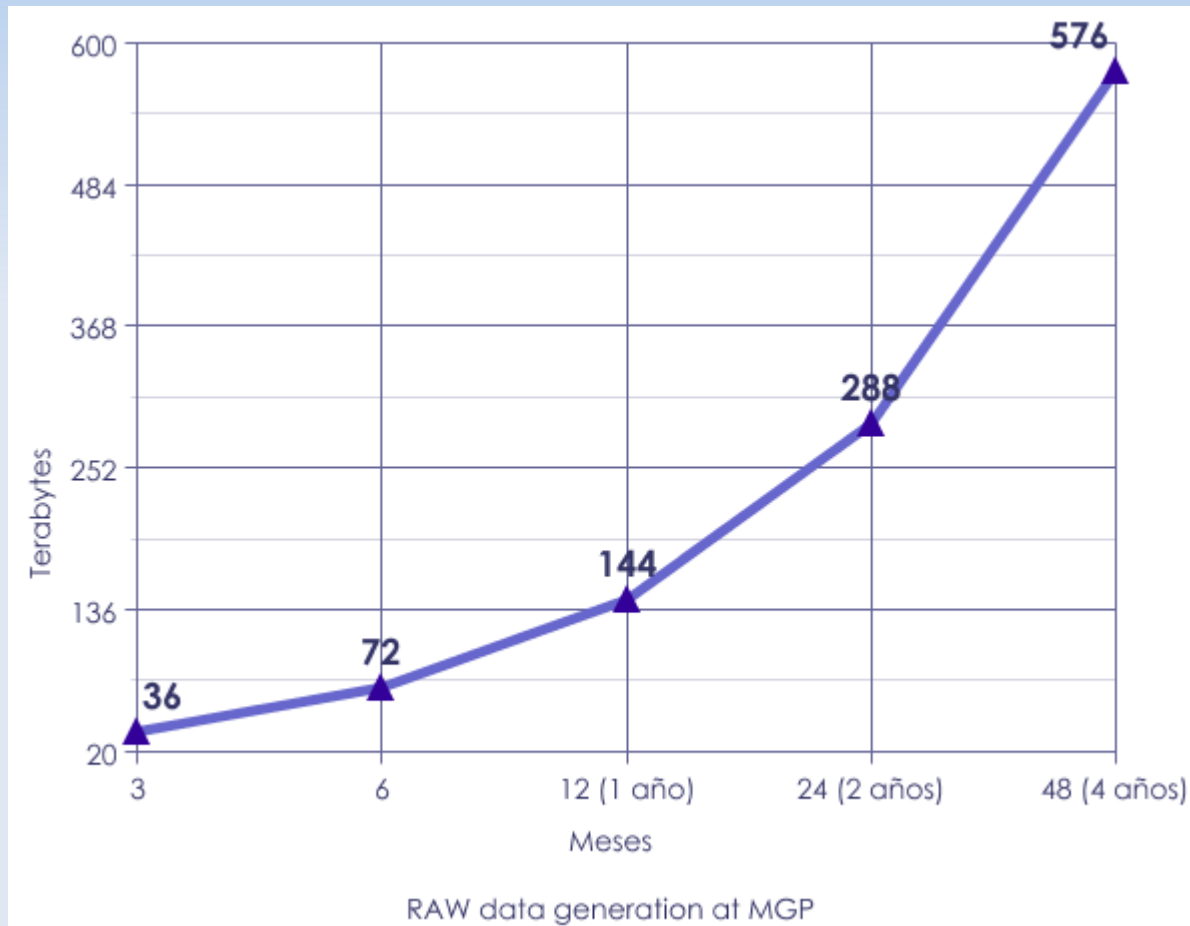


MGP raw data generation

- a solid sequencer run
 - 7 days running
 - Generates around 4TB
- Only the four solid sequencers working full time can generate around 12TB each week.
- 12TB just of raw data. After running bioinformatics analysis more data is generated
- Raw data size grows really fast
 - New sequencer models
 - New reagents



MGP raw data generation



Sequencing centers in Spain

CNAG

- Sequencing Instruments
 - 10 Illumina HiSeq2000
- Informatics infrastructure
 - 850 core cluster
 - 1.2 petabyte lustre filesystem (growing to 2PB)
 - 10 x 10 Gb/s link with marenostrom (Barcelona Super Computer 10,240 cores)

CNAG

Informatics



↑↓
10 x 10 Gb/s



Barcelona Super Computer 10,240 cores

850 core cluster supercomputer
1.2 petabyte harddiscs

BGI - Largest sequencing center in the world

- Sequencing Instruments
 - Illumina HiSeq
 - AB SOLiD System
 - Ion Torrent
- Informatics infrastructure (8 datacenters)
 - 20,576 cores cluster
 - 17PB

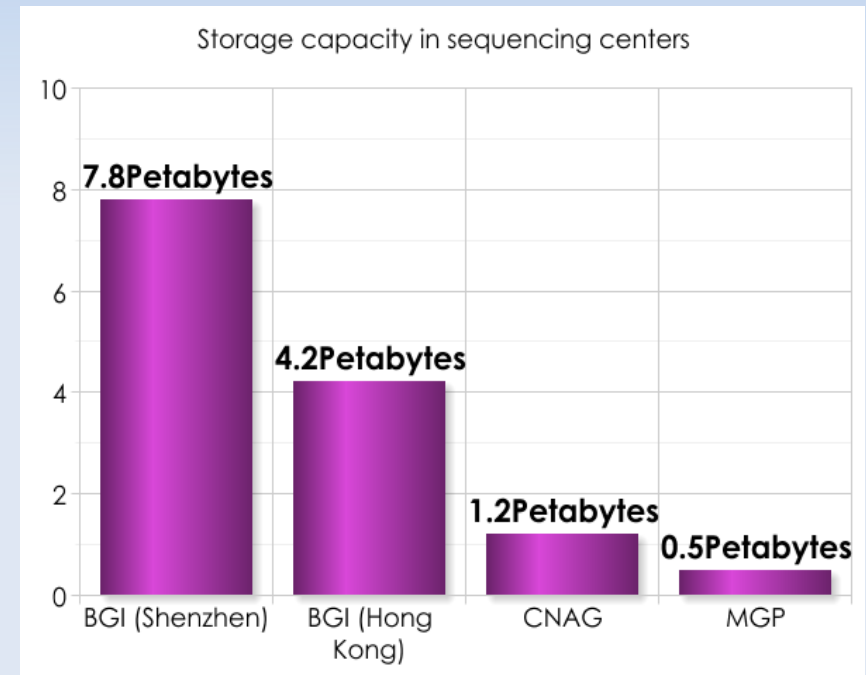
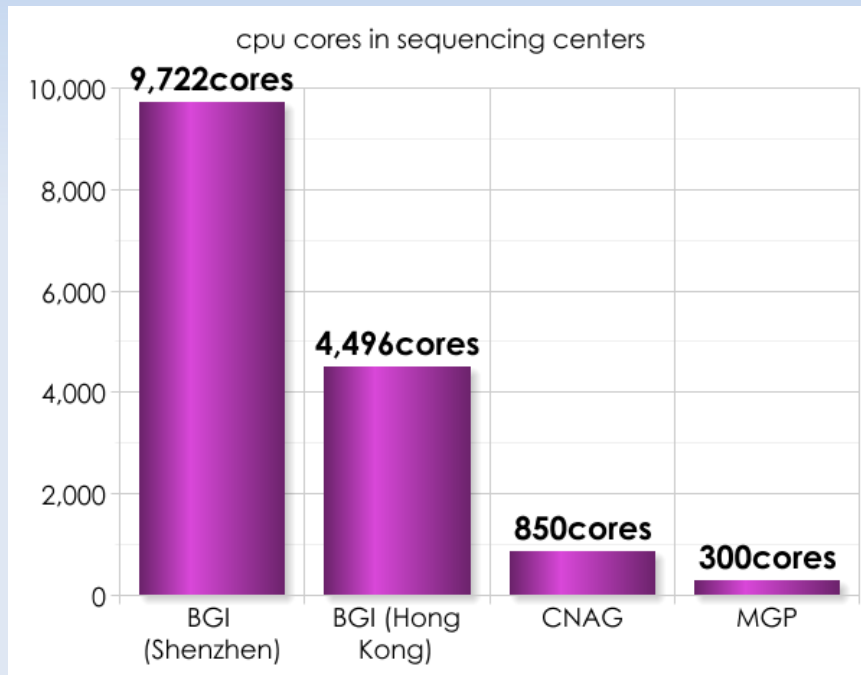
Source: http://www.genomics.cn/en/navigation/show_navigation?nid=4109

Largest sequencing center in the world

- Beijing Genomics Institute (BGI)



Sequencing center resources



Most used operating system is GNU/LINUX

TOP500 Statistics

TOP500 Release:

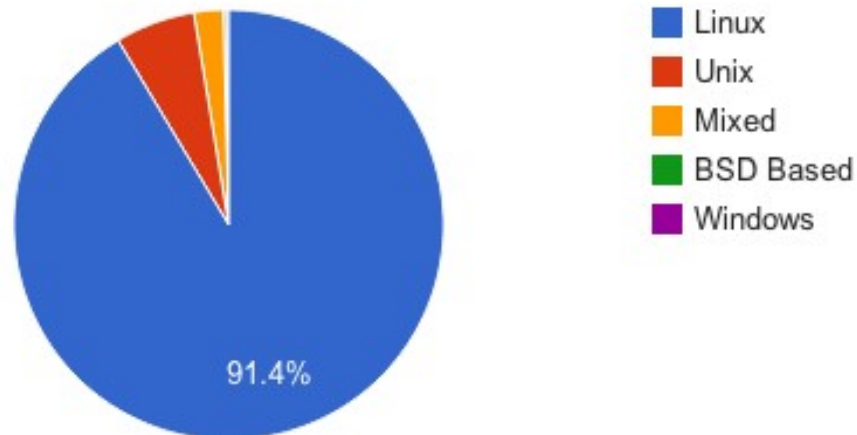
November 2011 ↕

Category:

Operating system Family ↕

R_{max} and R_{peak} values are in GFlops. For more details about other fields, check the [TOP500](#)

Operating system Family System Share



Source:

<http://www.top500.org/stats/list/36/osfam>

Alternatives – cloud computing

- Pros

- Flexibility.
- You pay what you use.
- Don't need to maintain a data center.

- Cons

- Transfer big datasets over internet is slow.
- You pay for consumed bandwidth. That is a problem with big datasets.
- Lower performance, specially in disk read/write.
- Privacy/security concerns.
- More expensive for big and long term projects.

