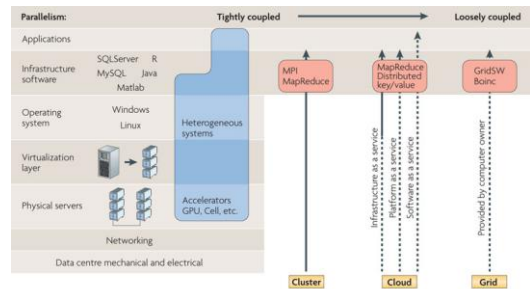


Infrastructure organization

BINF 630: Bioinformatics Methods

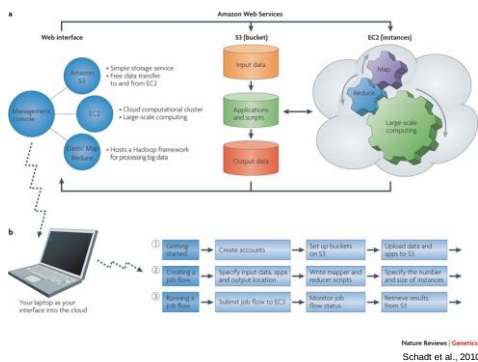
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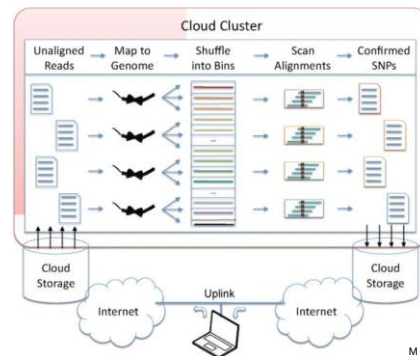
Nature Reviews | Genetics
Schadt et al., 2010

Cloud computing



Nature Reviews | Genetics
Schadt et al., 2010

Cloud computing and DNA sequencing



M C Schatz et al., 2010

Types of computational environments

	Computing architectures	Advantages	Disadvantages	Example applications
Large-scale computing platform				
Cluster computing	Multiple computers linked together, typically through a fast local area network, that effectively function as a single computer.	Cost-effective way to realize supercomputer performance.	Requires a dedicated, specialized facility, hardware, system administration and IT support.	• BLAST • Bayesian network reconstruction • Computing genetic associations in large-scale GWAS studies
Cloud computing	Computing capability that abstracts the underlying hardware architectures (for example, servers, storage and networking), enabling convenient, on-demand network access to a shared pool of computing resources that can be easily provisioned and released (NIST Technical Report).	The virtualization technology used results in extreme flexibility: good for one-off HPC tasks, for which persistent resources are not necessary.	Privacy concerns; less control over processes; bandwidth is limited as large data sets need to be moved to the cloud before processing.	• Searching sequence databases • Aligning new sequencing reads to genomes • General-purpose genomics tools (for example, GeneSifter from Geopit) • Most applications running on a cluster can be transferred to a cloud.
Grid computing	A combination of loosely coupled networked computers from different administrative centres that work together on common computational tasks, typically by volunteer computing efforts (such as Folding@home), which 'scrape' spare computational cycles from volunteers' computers.	Ability to enlist larger-scale computational resources at low or no cost (large-scale volunteer-based efforts).	By data transfers are difficult or impossible; minimal control over underlying hardware, including availability of cloud-based services.	• Protein folding (Folding@home) • Proteome analysis • Protein-protein interaction (HotSpot@home) • Predicting interactions between small molecules and proteins (Fold@home)
Heterogeneous computing	Computers that integrate specialized accelerators—for example, GPUs or reconfigurable logic (FPGAs)—alongside CPUs.	Cluster-scale computing for a fraction of the cost; optimized for computationally intensive fine-grained (parallel) local control of data and processes.	Significant expertise and programmer time required to implement applications; not generally available in cluster and cloud-based services.	• Bayesian network learning • Protein folding (Folding@home) • Molecular dynamics simulation (NAMD) • BLAST • CLUSTALW • HMMER • Reconstruction of evolutionary trees

The above categories are not exclusive. For example, heterogeneous computers are often used as the building blocks of clusters, grid or cloud computing systems; the shared computational clusters available in some organizations could be described as private Platforms as a Service (PaaS) clouds. The main differences between the platforms are degrees of coupling and memory—grid and cloud computers are designed for loosely coupled parallel workloads, with the grid resources allocated exclusively for a single time, whereas the underlying hardware resources in the cloud are typically shared among many users (multi-tenancy). Cluster computers are typically used for tightly coupled workloads and are often allocated to a single user (FPGA, field-programmable gate array; GPU, general-purpose processor); GPUs, graphics processing unit; GWA, genome-wide association; HPC, high-performance computing; NIST, National Institute of Standards and Technology.

Schadt et al., 2010

Types of computational environments

Environment	URL
Cloud computing	
Amazon Elastic Compute Cloud	http://aws.amazon.com/ec2
Bionimbus	http://www.bionimbus.org
NSF CluE	http://www.nsf.gov/cise/clue/index.jsp
Rackspace	http://www.rackspacecloud.com
Science Clouds	http://www.scienceclouds.org
Heterogeneous computing	
NVIDIA GPUs	http://www.nvidia.com
AMD/ATI GPUs	http://www.amd.com
Heterogeneous cloud computing	
SGI Cyclone Cloud	http://www.sgi.com/products/hpc_cloud/cyclone
Penguin Computing On Demand	http://www.penguincomputing.com/POD/Summary

GPU, graphics processing unit; NSF, US National Science Foundation.

Schadt et al., 2010

Defining Big Data

NOT JUST SIZE

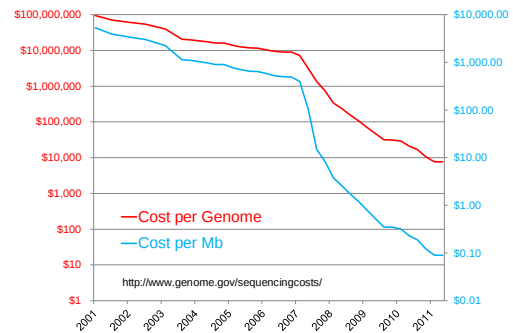
The three Vs of Big Data: volume, variety and velocity
(D.Laney, 2001)

Elements of "Big Data" include:

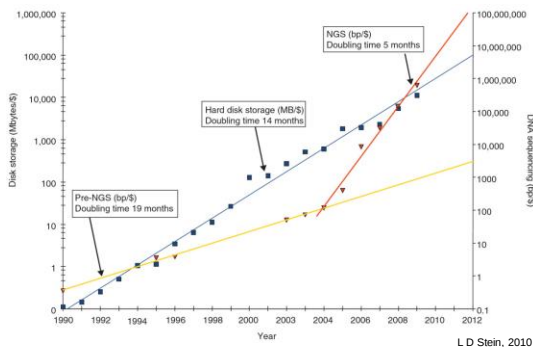
- The degree of complexity within the data set
- The amount of value that can be derived from innovative vs. non-innovative analysis techniques
- The use of longitudinal information supplements the analysis

http://mike2.openmethodology.org/wiki/Big_Data_Definition

Genome sequencing costs



Sequencing and storage cost



Data Mining

- Data mining is the exploration and analysis, by automatic or semiautomatic means, of large quantities of data in order to discover meaningful patterns and rules
- Common data mining tasks
 - Classification
 - Estimation
 - Prediction
 - Affinity Grouping
 - Clustering
 - Description

Knowledge Discovery

Knowledge is a pattern that exceeds certain threshold of interestingness.

Factors that contribute to interestingness:

coverage
confidence
statistical significance
simplicity
unexpectedness
actionability

Knowledge Discovery

- Directed and Undirected KD
- Directed KD
 - Purpose: Explain value of some field in terms of all the others
 - Method: We select the target field based on some hypothesis about the data. We ask the algorithm to tell us how to predict or classify it
 - Similar to hypothesis testing (e.g., in regression modeling) in statistics

Knowledge Discovery

- Undirected KD
 - Purpose: Find patterns in the data that may be interesting
 - Method: clustering, affinity grouping
 - Closest to ideas of machine learning in artificial intelligence
- Comparison
 - UKD helps us to recognize relationships & DKD helps us to explain them

Classification

- Classifying observations into different categories given characteristics

Estimation

- Rules that explain how to estimate a value given characteristics

Prediction

- Rules that explain how to predict a future value or classification, given characteristics

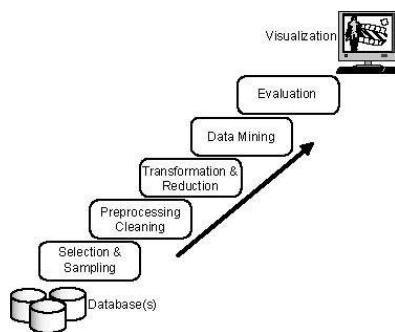
Affinity Grouping

- Grouping by relations (not by characteristics)

Clustering

- Segmenting a diverse population into more similar groups
- In clustering, there are no pre-defined classes and no examples. Records are grouped together by some similarity measure.

Knowledge Discovery



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