



Genome Data Management Workshop

January 22nd-26th 2011

Universidad de Puerto Rico en
Mayagüez

GDM Workshop

Day 1

- I. DNA sequencing
- II. Planning your project
- III. Hardware requirements
- IV. A primer on Linux

Day 2

- V. Genome assembly

Day 3

- VI. Genome annotation

Day 4

- VII. Perl
- VIII. Phylogeny

Day 5

- VIII. Phylogeny

WIFI Network code
SquawkinGood

II. Planning your project

3 questions

What is your goal?

What is the best way to achieve it?

How much can you spend?

Evaluate your strengths

Outsourcing

Got DNA/RNA?

Libraries?

Sequencing?

Bioinformatics?

A collaboration?

Additional costs

PCRs

Cloning

Sanger sequencing

Misc. lab costs

III. Hardware requirements

The hidden cost

1 Gigabase of DNA sequence \approx 1 GB of RAM

32 bits vs 64 bits vs 128 bits

32 bits = 2^{32} = 4 294 967 296 = 4 Gigabytes of RAM

64 bits = 2^{64} = 18 446 744 073 709 551 616 = 16 Exabytes of RAM

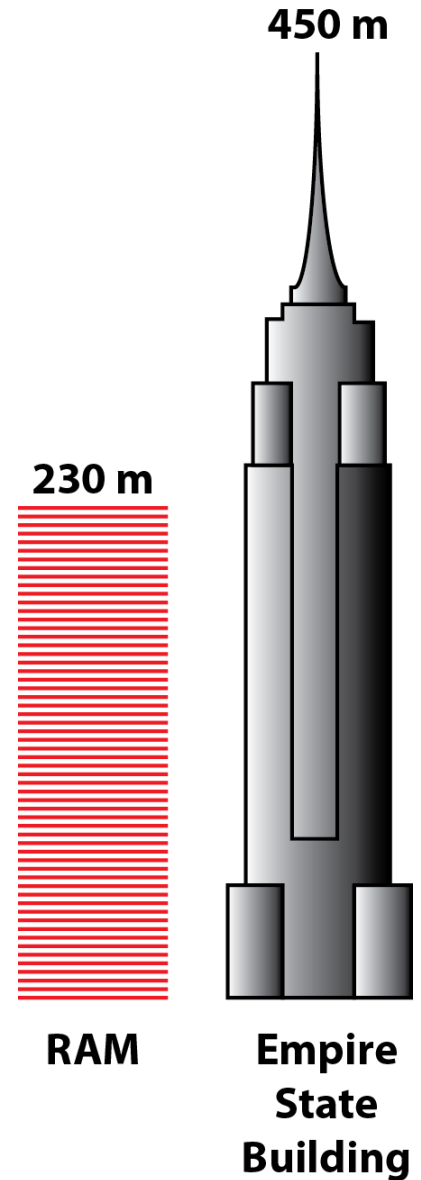
128 bits = 2^{128} = do we even have a number for that? ^ Not quite

256 TB of RAM

256 TB = 262 144 GB

At current density (8GB per chip) = 32 768 chips

At a thickness of $\approx 7\text{mm}$, if piled up...



Operating systems (64 bits)

MS Windows 7

Starter	2 GB
Home Basic	8 GB
Home Premium	16 GB
Professional, Enterprise, Ultimate	192 GB

Windows Server 2008 R2

Datacenter, Enterprise	2 TB
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Mac OSX

Leopard/Snow Leopard 32/64 bits kernels	unspecified
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Linux

44 bits kernel	16 TB
45 bits kernel	32 TB
46 bits kernel	64 TB

Disk space

Data files	1-100 Gigabytes
Temporary files	can easily range in Terabytes
Current SATA drives	3 Terabytes

"Let's face it, we're not changing the world. We're building a product that helps people buy more crap - and watch porn."

Bill Watkins, Seagate CEO - 2006

I/O speed

Optical media

Not a viable option

SSD

Fast, small, expensive

Hard drive

Slower, bigger, cheaper

RAID*

Software or hardware

*Not the WoW kind...

File compression

Saves on disk space

Faster I/O

Requires more RAM & CPU (at launch)

Works well on text files (*i.e.* most sequencing output files)

The beasts of burden

Supercomputers

Local clusters

Workstations

Mobile workstations

Cloud-based, networked or local?



Computing resources



Compute/Calcul Canada
<https://computecanada.org/>



TeraGrid
<https://www.teragrid.org/>

A supercomputer (Le colosse)



CPU	960 nodes Xeon X5560 @ 2.8GHz (7680 cores)
RAM	24GB/node (23TB total)
Disk	500TB Lustre FS



Supercomputer/Clusters limitations

RAM per node

Disk quota

Bandwidth

Queuing & CPU time limit

User permissions

IBM BladeCenter HX5 Express MAX5

CPU 2x Intel Xeon X6550 Quad Core 2.0GHz

RAM **112GB** DDR3 ECC

22 800\$ + tax

IBM BladeCenter HX5 Express MAX5

CPU 2x Intel Xeon E7540 Quad Core 2.0GHz

RAM **64GB** DDR3 ECC

15 000\$ + tax

My own rig (custom built)

CPU 2x Intel Xeon E5506 Quad Core 2.13GHz

RAM **96GB** DDR3 ECC

5 500\$ + tax

Backups

DVDs

Too small

Blu-Rays

Might do it

External disks

Decent option

Tape drives

Decent option

NAS

Better option

IV. A primer on Linux

What is Linux?



Free operating system created by Linus Torvalds & Richard Stallman

Similar to UNIX by AT&T (Bell Labs)

Actively developed under the GNU General Public License

Source code is freely available

The Linux distributions



Fedora

<http://fedoraproject.org/>



Ubuntu

<http://www.ubuntu.com/>



Red Hat

<http://www.redhat.com/>



openSUSE

<http://www.opensuse.org/>



Gentoo

<http://www.gentoo.org/>

And many more...

How?



From a live CD/DVD or install CD/DVD



From within Windows (WUBI)



From a flash drive



From the web via HTTP or FTP

The Linux kernel

The command shell

BASH (Bourne-Again SHell)

TCSH

DASH (Debian Almquist SHell)

KSH

The X interface

The window managers



IceWM

<http://www.icewm.org/>



Enlightenment

<http://www.enlightenment.org/>



Sawfish

<http://sawfish.wikia.com/>

And many more...



Fluxbox

<http://fluxbox.org/>

The desktop managers



Gnome

<http://www.gnome.org/>



KDE

<http://www.kde.org/>



XFCE

<http://www.xfce.org/>



LXDE

<http://lxde.org/>

Compilers/interpreters

Source code

Programming languages

The compilers

Binaries

The partitions

Boot

Swap

Root (/)

The user types

Root

Sudoers

Normal users

Groups

The user permissions

The owner d**rw**xrwxrwx

The group drwx**rw**xrwx

The others drwxrwx**rw**x

r = read, w = write, x = execute, - = permission denied

Basic commands

cd	change directory
cp	copy files/directories
mv	move/rename files/directories
ls	list the content of a directory
rm	delete files/directories
*	the wildcard
>	redirect the bash output to a file
>>	append the bash output to a file

Useful commands

pwd	display your current directory
tar	compress files/folders
ln	create aliases (links)
top	show processes
jobs	display current jobs
kill	terminate jobs/processes
screen	detachable Shell
shutdown	stop/reboot computer

Useful tips

Tab-typing

cd \$HOME

cd ~

Ctrl+C

history

up and down arrows

less/more

tail

autocomplete your command

go back to your home directory

another way to go back to \$HOME

abort a process

show your last commands

scroll through previous commands

reads the 1st lines of a text file

reads the last lines of a text file

Running the analyses

Command lines vs. GUI

Locally

Remotely

SSH & SFTP

Installing additional software

The easy way

The hard way

The Path

A word on dependencies

Need help?

<http://embnet.org/en/QuickGuides>