



Computational Cluster

User Cheatsheet - R1.3 - 05/12/19

Connecting with Kabré

Establishing an SSH session	Open a terminal program and type: \$ ssh user@kabre.cenat.ac.cr Then press enter three times when requesting the creation of SSH keys. Exit session: \$ exit
	In your computer, open a terminal and type: \$ ssh-keygen -t rsa -C "your_email@example.com" \$ ssh-copy-id user@kabre.cenat.ac.cr
Copying files	From your computer to Kabré: \$ scp files [user]@kabre.cenat.ac.cr:[path] From Kabré to your computer: \$ scp [user]@kabre.cenat.ac.cr:[files] [path] These commands must be executed in your computer. \$ scp -r user@host:[directory] [directory] Copy all files and directories recursively.
	\$ ssh user@kabre.cenat.ac.cr \$ passwd

Kabré's Queues System

Writing a SLURM job file	#SBATCH -job-name=<job_name> #SBATCH -output=<result_name> #SBATCH -partition=<partition_name> #SBATCH -ntasks=<multiply X*Y> #SBATCH -time=<HH:MM:SS> execute your program here
	\$ sbatch job.slurm
Submitting your job	
Monitoring your jobs	In Kabré, type: \$ watch -n 5 squeue
Retrieving results	All jobs will generate an output file: <result_name>
Interactive jobs	\$ salloc

Environment Modules

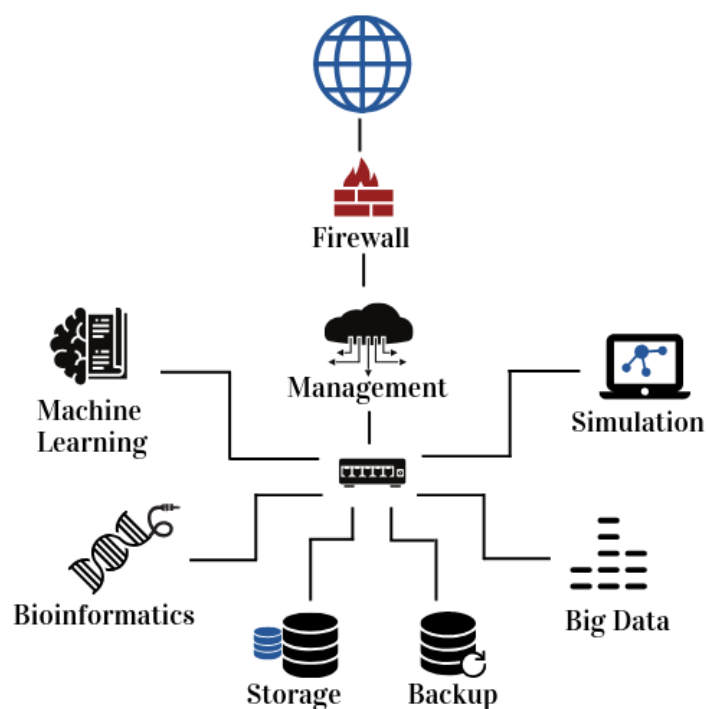
List loaded modules	\$ module list
List available modules	\$ module avail
Load a module	\$ module load module_name
Unload a module	\$ module unload module_name



Centro Nacional de Alta Tecnología
Edificio Dr. Franklin Chang Díaz, Pavas.
www.kabre.cenat.ac.cr
cnca@cenat.ac.cr
Tel: (506) 2519 5839



Kabré's composition



Virtualization

Just don't mess up here!

Don't execute programs here! These nodes are a shared-working area, use them to:

- ▶ Create and edit files
- ▶ Create directories and move files
- ▶ Copy files to and from your computer
- ▶ Compile code
- ▶ Submit jobs
- ▶ Manage your active jobs

Login-nodes

Nu (Simulation)

Each blade has 4 Intel Xeon Phi KNL nodes with 64 cores @ 1.3 GHz and 96 GB

Nukwa (Machine Learning)

Nvidia Tesla K40. Hoster has an Intel Xeon with 4 cores @ 3.2 GHz and 16 GB

Andalan (Big Data)

Each blade has 2 Intel Xeon E5-2650 v4 with 12 cores @ 2.20 GHz and 64 GB

Dribe (Bioinformatics)

-

Available Queues

Name	Platform	Number of nodes	Time slot (in hours)
nu	Xeon Phi KNL	1	72
nu-debug	Xeon Phi KNL	1	8
nu-wide	Xeon Phi KNL	12	24
nu-long	Xeon Phi KNL	1	744
nukwa	GPU	1	72
nukwa-debug	GPU	1	8
nukwa-wide	GPU	2	24
nukwa-long	GPU	1	168