



# Introduction to the OMNI cluster

Jan Steiner Zentrum für Informations- und Medientechnik

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#### A word about Zoom

- Exercises:
  - -Groups of three
  - -One person shares screen
  - -Solve cooperatively
  - -Screen-sharer switches for next exercise
  - -I will visit each group





# Who am I

- Jan Steiner
  - -Aerospace Engineering, Uni Stuttgart (grad. 2010)
  - -German Aerospace Center Braunschweig (fluid dynamics)
  - -At ZIMT since July 2017
- Area (with one other colleague):
  - -HPC training and support
  - -Training courses (once every semester)
    - This course
    - Linux
- Additional support: performance optimization





### **Round of introductions**

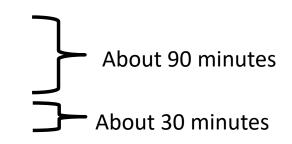
- What department/institute are you with?
- What is your field / research topic?
- How do you use / intend to use the cluster?
- What is your previous experience?
- Is there something specific you want to learn today?





# Outline

- 1. Getting onto the cluster
  - Structure of a cluster
  - Getting access and help
  - Connecting to the cluster
  - Exercise 1: setup, login
- 2. Using the cluster
  - Workspaces
  - Environment modules
  - Jobs
  - Exercise 2: your first job script
- 3. SLURM explained
  - Tasks, processes, cores
  - Miscellaneous SLURM stuff
  - Exercise 3: SLURM options









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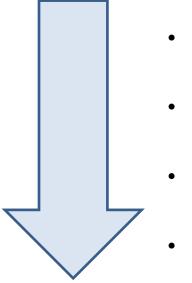
# Background

- Computations can become too large for one computer
  - Too much concurrent data for RAM
  - Too much total data for hard drive
  - -Execution time in months, years or more
  - Too many small problems (e.g. parameter study)
- $\rightarrow$  Use more computers
- <u>Cluster</u> of computers
  - Components similar to PC
  - But many, and interconnected





### Physical structure of a cluster



- Core (Processor)
- Node (Blade)
  - Rack (Cabinet, Chassis)
- Cluster (Supercomputer)





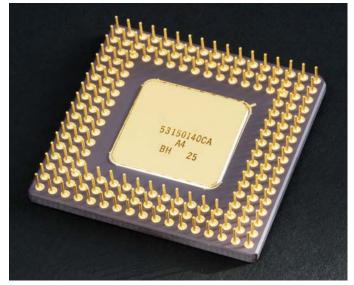






#### Cores

- (Almost) identical to PC processors
- General purpose
- Hyperthreading (two cores in one): disabled on clusters
- Sometimes specialized
  - E.g. graphics processors (GPU)
  - Limited operations, but faster



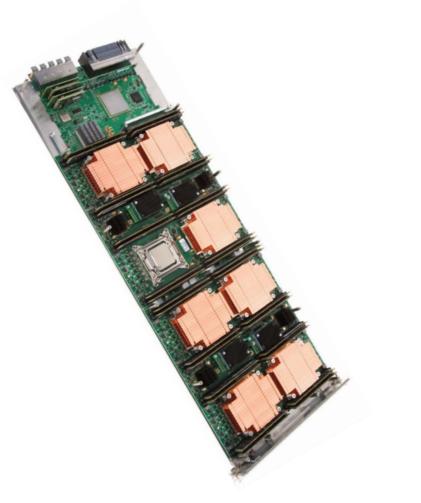
Source: Wikimedia Commons





# Nodes

- Similar to PC motherboards
- 2-4 CPUs, each with many cores
- Usually central RAM – OMNI: 256 GB
- Types
  - -Compute, Login, Management
  - "Fat" (more RAM), GPU
    - smp1: 1536 GB RAM







# Cabinet

- Houses multiple nodes
- Cooling
- Power supply
- Interconnect (Network)
  - -Faster than regular Ethernet
  - -Makes cluster a cluster
  - -OMNI: Infiniband







# Cluster

- Multiple cabinets
  - -OMNI: 9 cabinets, ~550 nodes, 29000 cores
- Infrastructure (e.g. fire suppression)
- Central file storage (hard disks)
  - Sometimes individual nodes have hard disks







# Situation at Uni Siegen

- Current: multiple systems
  - -OMNI cluster
  - HPE Moonshot (HTC nodes)
  - -NEC Aurora vector computer
  - -ARM cluster

- Main cluster: OMNI
  - -Since early 2021
  - -3-4 times more regular CPUs than previous HoRUS cluster
  - -nVIDIA GPUs





### **OMNI cluster hardware**

- 434 regular compute nodes
  - -hpc-node001-hpc-node136
  - -2x32 AMD EPYC Rome CPUs, 256 GB RAM each
- 2 SMP (Shared Multiprocessing) nodes
  - -smp-node001/002
  - -64 CPUs, 1536 GB RAM each
- 10 GPU nodes with total of 24 GPUs
  - -gpu-node001-010
  - -NVIDIA Tesla V100
  - -4x1, 2x2, 4x4 GPUs





Sources: Wikimedia Commons





# **OMNI cluster hardware**

- 4 login nodes hpc-login01-04
  - Identical to compute nodes except
     512 GB RAM
- 2 management nodes (not accessible to users)
- Around 500 TB total hard drive space

   Additionally 32 TB RAM SSDs ("Burst Buffer")
- Various network components etc...







# Situation at Uni Siegen

- HPE Moonshot HTC System
  - -45 nodes (2x login, rest compute)
    - 8 CPUs, 64 GB RAM each
  - -Designations: htc001htc007
  - -Shares homes with OMNI
  - -High-Throughput Computing:
    - Smaller jobs, but more



Source: hpe.com





# Situation at Uni Siegen

- NEC SX Aurora Tsubasa System
  - -2 machines ("vector host")
    - 2 cards ("vector engines") each
  - Intended for testing vector architecture
    - Similar to GPUs
  - -Names: vec01-vec02

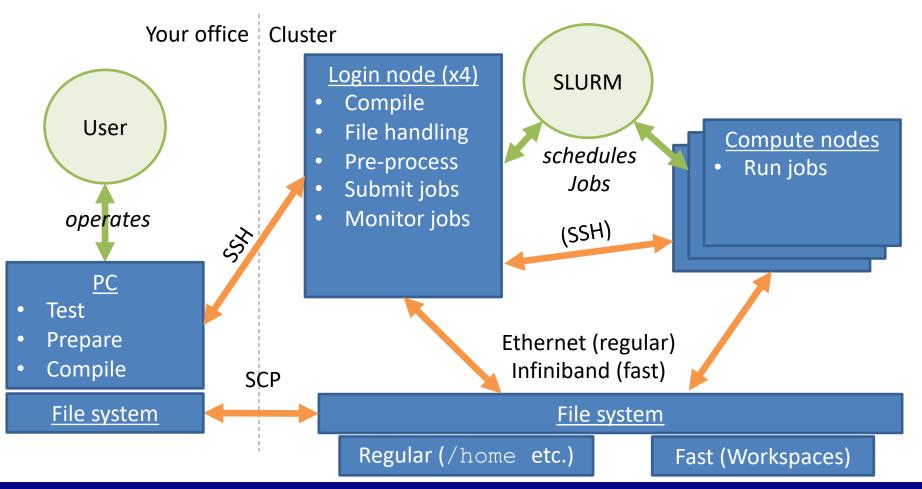


Source: nec.com





# Logical structure of a cluster







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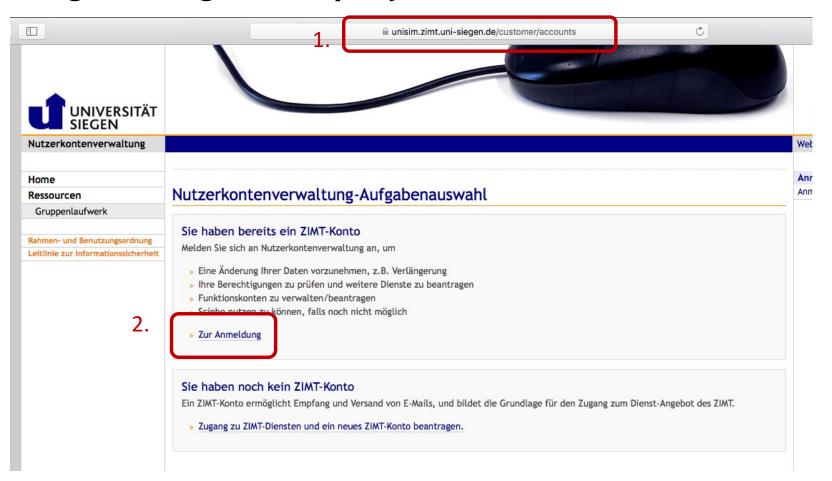
#### Getting cluster access

- Register for cluster access
  - Employees: Nutzerkontenverwaltung
  - -Students: need an employee supervisor
  - -Valid for all our systems
- Set up an SSH connection
  - -Explained in a moment





#### **Registering an employee**







#### Registering an employee

Nutzerkontenverwaltung des ZIMT an der Universität Siegen.

B	Benutzerkennung: js05	56352	
P	Passwort:	••••••• •••	
	Log	g In	
		ITÄT	Z(MT)
	Nutzerkontenverwalt		Webmail unisono Formular
			Annual Hermite
	Home	Noire Optioner	Anmeldung Welcome is056352
	Ressourcen	Meine Optionen	Welcome js056352
	Ressourcen Meine Anträge		_
	Ressourcen	Meine Optionen         Für mich selbst buchbare Optionen	Welcome js056352 Gültigkeit: unbefristet Abmelden
	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste	Für mich selbst buchbare Optionen	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte
I. (	Ressourcen Meine Anträge		Welcome js056352         Gültigkeit: unbefristet         Abmelden         Ich möchte         meine Nutzungsdauer verlängen
4. <b>(</b>	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste	Für mich selbst buchbare Optionen	Welcome js056352         Gültigkeit: unbefristet         Abmelden         Ich möchte         meine Nutzungsdauer verlängen
4. <b>(</b>	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste Meine Optionen Meine Groppen Passwortänderung	Für mich selbst buchbare Optionen           Ansys         Gebucht         Keine nachträgliche Änderung möglich         Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte
4. <b>(</b>	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste Meine Optionen Meine Sroppen	Für mich selbst buchbare Optionen           Ansys         Gebucht         Keine nachträgliche Änderung möglich         Hilfe	Welcome js056352         Gültigkeit: unbefristet         Abmelden         Ich möchte         meine Nutzungsdauer verlängen
I. (	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste Meine Optionen Meine Groppen Passwortänderung Helpdesk	Für mich selbst buchbare Optionen         Ansys       Gebucht       Keine nachträgliche Änderung möglich       Hilfe         MAXQDA Campus Lizenz       Nicht gebucht       Jetzt buchen       Hilfe         Sophos Antivirus       Nicht gebucht       Jetzt buchen       Hilfe	Welcome js056352         Gültigkeit: unbefristet         Abmelden         Ich möchte         meine Nutzungsdauer verlängen
. (	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste Meine Optionen Meine Groppen Passwortänderung Helpdesk Suchen	Für mich selbst buchbare Optionen         Ansys       Gebucht       Keine nachträgliche Änderung möglich       Hilfe         MAXQDA Campus Lizenz       Nicht gebucht       Jetzt buchen       Hilfe         Sophos Antivirus       Nicht gebucht       Jetzt buchen       Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte meine Nutzungsdauer verlänger weitere ZIMT-Dienste beantrage
. (	Ressourcen Meine Anträge Meine Daten Meine Konten/Dienste Meine Optionen Meine Grappen Passwortänderung Helpdesk Suchen Bestandskontenübern.	Für mich selbst buchbare Optionen         Ansys       Gebucht       Keine nachträgliche Änderung möglich       Hilfe         MAXQDA Campus Lizenz       Nicht gebucht       Jetzt buchen       Sophos Antivirus         Sophos Antivirus       Nicht gebucht       Jetzt buchen       Sophos Antivirus	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte meine Nutzungsdauer verlänger weitere ZIMT-Dienste beantrage





W

#### Registering an employee

#### Meine Optionen

für mich se	lbst buchbare Optionen			
	Ansys	Gebucht	Keine nachträgliche Änderung möglich	Hilfe
	MAXQDA Campus Lizenz	Nicht gebucht	Jetzt buchen	
	Sophos Antivirus	Nicht gebucht	Jetzt buchen	
	Microsoft 365	Gebucht	Keine nachträgliche Änderung möglich	Hilfe
	Stata Campus Lizenz	Nicht gebucht	Jetzt buchen	
	Adobe Acrobat Pro DC	Gebucht	Buchung ändern	
	Adobe VIP-Vertrag	Nicht gebucht	Jetzt buchen	
	Code42 Clientbackup Angebot der GWDG	Gebucht	Keine nachträgliche Änderung möglich	Hilfe
5.	Ressourcen zum Wissenschaftlichen Rechnen (OMNI)	Gebucht	Keine nachträgliche Änderung möglich » OMNI » Horus » HPE Moonshot	
			Hilfe	

#### Introduction to the OMNI cluster





#### **Registering an employee**

#### 6.

#### So geht es weiter

- » Drucken Sie den Antrag jetzt aus:
- Drucken
- (öffnet ein neues Fenster oder einen neuen Reiter)

- » Unterschreiben Sie den Ausdruck!
- » Lassen Sie den Ausdruck durch den angegebenen Vorgesetzten unterzeichnen!
- » Lassen Sie den Dienststempel der Einrichtung stempeln.
- » Hinweis: Ohne Unterschriften und Dienststempel wird der Antrag abgelehnt!
- » Senden Sie den Ausdruck an den ZIMT-Benutzerservice; der Ausdruck verfügt über die Adresse
- » Sollte nach vier Wochen der unterschriebene und gestempelte Antrag nicht beim Benutzerservice eingegangen sein, wird der Vorgang ohne weitere Benachrichtigung gelöscht.





- Every student account has an assigned supervisor
  - Supervisor adds student in Nutzerkontenverwaltung
  - -Supervisor is responsible
- Student accounts time out after 1 year
  - -Can be extended by supervisor
  - -Warning before time-out, data not immediately lost
- Otherwise, no restrictions for student accounts





Nutzerkontenverwaltung des ZIMT an der Universität Siegen.
--

Benutzerkennung: js05635						
asswort:		<b>9</b> ~				
	Log In					
	/ERSITÄT					ZMT
SIEGI Nutzerkontenve	EN					Webmail unisono Formular
Home						Anmeldung
Home Ressourcen		Meine Optionen				Anmeldung Welcome js056352
		Meine Optionen				Welcome js056352 Gültigkeit: unbefristet
Ressourcen						Welcome js056352
Ressourcen Meine Anträge	iensie	Meine Optionen Für mich selbst buchbare Optionen				Welcome js056352 Gültigkeit: unbefristet
Ressourcen Meine Anträge Meine Daten			Gebucht	Keine nachträgliche Änderung möglich	Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte
Ressourcen Meine Anträge Meine Daten Meine Konten/D		Für mich selbst buchbare Optionen	Gebucht	Keine nachträgliche Änderung möglich	Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte meine Nutzungsdauer verlängen
Ressourcen Meine Anträge Meine Daten Meine Konten/D Meine Optionen		Für mich selbst buchbare Optionen	Gebucht Nicht gebucht	Keine nachträgliche Änderung möglich Jetzt buchen	Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden
Ressourcen Meine Anträge Meine Daten Meine Konten/D Meine Optionen Meine Gruppen		Für mich selbst buchbare Optionen Ansys MAXQDA Campus Lizenz	Nicht gebucht	Jetzt buchen	Hilfe	Welcome js056352 Gültigkeit: unbefristet Abmelden Ich möchte meine Nutzungsdauer verlängen
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Testbetrieb nur ZIMT experimentell	Sciebo - die Campuscloud	Nicht gebucht	Jetzt buchen
Testbetrieb nur ZIMT experimentell	GitLab der Universität Siegen	Gebucht	Buchung ändern
reigabe vor	n ggf. kostenpflichtigen Optionen für D	ritte	
Adobe Acrobat	Pro DC für Studenten/Mitarbeiter	Zur	r Eingabe
Adobe VIP-Vert	rag für Studenten/Mitarbeiter	Zur	r Eingabe





Zusammenstellen von Empfängern: Ressourcen zum Wissenschaftlichen Rechnen (OMNI)

<b>Bezug</b> Bitte geben Sie die die freizugebenden		Check this mark haftlichen Rechnen (OMNI) für die gewünschten Personen ein und wählen Sie		
E-Mail-Adresse		Ressourzen zum Wissenschaftlichen Rechnen (OMNI)		
		+		
Abbrechen	Must be Uni Siegen address	Weiter		
	Add multiple	people in one go if desired		





# After registration

- Prompt to agree to Terms of Use
   Obstacle for people from some countries (Iran)
- Account not immediately ready
  - -Usually next day
  - -Contact us if still no e-mail after a week
- When ready: "Welcome to the OMNI cluster" e-mail
  - -Keep this e-mail, it contains the cluster address(es)

Demo 1





# Getting help

- Cluster website: <u>https://cluster.uni-siegen.de</u>
  - -Usage information (like our courses)
  - -What is installed
- Consult documentation, internet

   Built-in help man <command> or <command> -h or --help
- Consultation hour (Zoom)
  - Every Tuesday 2 PM 3 PM
  - -Online (link on cluster website  $\rightarrow$  Events page)
- Support e-mail address: <u>hpc-support@uni-siegen.de</u>

Demo 2





### Problems

- Open a ticket
  - -Email to <u>hpc-support@uni-siegen.de</u>
  - -Centralized ZIMT ticket system
  - -Tell us what error (message) is
    - For jobs: attach job script, log file
- Please don't email us directly
  - -Person might be on vacation etc.
  - -Entire team has an overview what's wrong
  - -Also not good: <u>hpc-team@uni-siegen.de</u>





#### How to use other resources

- This course covers mostly OMNI cluster
- Using HPE Moonshot: relatively easy, similar to OMNI
- Other resources: get in contact with us





### Special cases

- Jupyter portal:
  - -Not yet ready!
  - -Enter Jupyter portal address in Browser
- Adding students for a teaching event
  - -Allowed in principle, contact us!
  - -Moonshot nodes intended for this purpose
  - -We may set up a reservation to avoid wait times (on a case-bycase basis)
- Absolutely not allowed: giving your password to another person





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#### Connecting to the cluster

- You can connect from any system via console
  - -Linux: Easiest
  - -Mac OS: Relatively easy
  - -Windows: now also built in
- Outside university network:
  - -Needs VPN for user/password access
  - -VPN not necessary for key-based access





# SSH Software

- Clusters typically accessed via Secure Shell (SSH) protocol
- Most commonly OpenSSH software
- Available for all operating systems
  - -Linux: original
  - -Mac OS: basically identical
  - -Windows 10 (since 2019): integrated in cmd/Powershell
- Additional tools, especially on Windows: Putty, MobaXTerm





## SSH Basic Use

- Connect with  $\mathtt{ssh}$  command

ssh [options] <username>@<hostname>

• You will be asked for password

-Alternative: set up public/private key pair (later)

- Can specify configurations to simplify login
- Console-based, but opening windows possible
- Multiple simultaneous connections possible

Demo 3





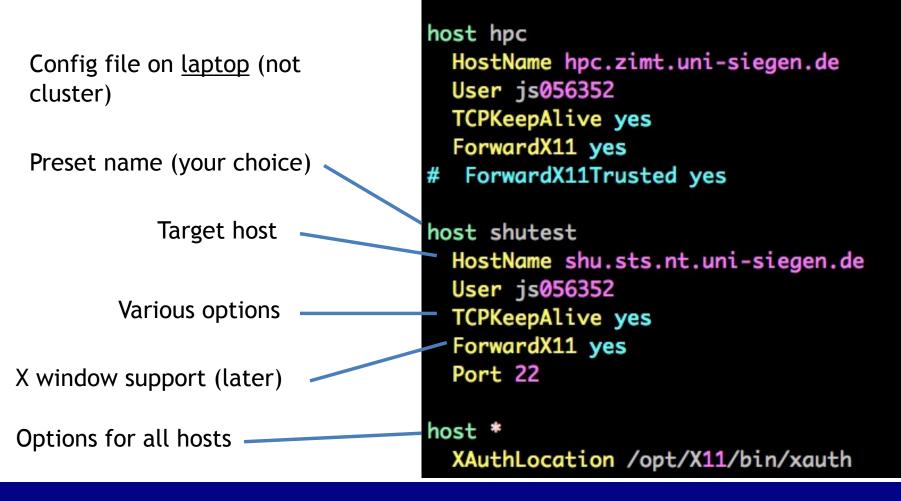
# **SSH Configuration**

- OpenSSH allows presets
- Can create text file ~/.ssh/config
  - -Edit if already exists
- One preset per connection (cluster etc.)
  - -Specify username
  - -Other options (many possibilities)
- Log in with ssh <presetname> instead of ssh [options] <user>@<host>





# SSH Configuration File







# SSH Key-based authentication

- Login with public/private key pair instead of password
- Convenient
  - -Good for automated connections
- Potentially more secure
- Only as secure as your PC
  - -Treat private key file like a physical key





# Key pair workflow

- 1. You generate key pair
  - On your PC
  - Tool ssh-keygen (comes with OpenSSH)
  - Keys are text files in ~/.ssh directory
- 2. You copy public key to cluster
  - ssh-copy-id (comes with OpenSSH)
  - Windows: manually copy and paste key
- 3. When logging in, OpenSSH will select key





### **Key generation**

- Run SSH key generator
  - 1. On local PC, type ssh-keygen
  - 2. Enter filename for new key
    - Should be inside ~/.ssh directory
    - **Caution:** will overwrite without asking
  - 3. Enter passphrase
  - 4. Confirm passphrase





### Copy key to cluster

- On <u>local</u> PC, use ssh-copy-id command
  - -Syntax: ssh-copy-id -i <keyfile> <user>@<host>
  - -Not available on Windows
  - -Remember you need to be inside Uni network/Uni VPN
- Alternative: copy manually
  - -On local PC, open public key file with text editor
  - -One line of text, three parts: algorithm, key, comment
  - -On <u>cluster</u>, open ~/.ssh/authorized\_keys
  - -Paste line, adjust comment as needed





# Key selection and tips

• When logging in, key will be used automatically

-May specify key file manually if needed (option -i)

- If you get asked for password, key not recognized
- Tips:
  - -Use one key per PC (in case of theft/compromise)
  - -Not recommended to leave passphrase empty
    - But only needs to be entered once

Demo 4





# Public key format

- One line per key (e.g. authorized\_keys)
- Three elements:
  - Encryption algorithm
  - Public key
  - -Comment
- Comment may be adjusted (from which device)





### Public key example

#### Algorithm

ssh-rsa

AAAAB3NzaC1yc2EAAAADAQABAAABgQC+iMPDyFgXxpIF8r0rJFFXY0S/Gy/l 1ijXuEs564y6cG/F95uxTjEC0kJEdmtix8lYfN8eIQ92xxib4/WQ0A243oh2 svD10R3gKNtzyjvc+eNMPOgf2yY0IlV1I3GfwGgLnTSKYgQH6HGcUgb/nQF3 eCcB9r3WCyoZ/tm4DrvuU8RQCD50fpq6D1p5017JaXCNSor9sbXqSSodnjTR nFyDSDf3sUGQuUUdXX1B1H35Kn2CWEP142ivTnehcUHDodRXmLIcrplId1F jDIEF1TP3h4FHeTcEY4ibHZzxgpbJ60By6mCq3AuNnriSu6adCRIBkZ81giR VpilSouzAbXQofzfJrzFnVcEDtPzbNUH6VCS32KMddlssraQmCwtEtFfu9nG C9F+dXoV38ZxQs9F4cUjqLGtkv51C0viGhadWmkpn0Ir5VdV7Vu319wWZ5wL FCrh/RR7S0UuIfr3VcnrK58F16yM1A/i8i3rQyAnZBn86/YwfnnIFRE9C5sA KAU= jansteiner@Jans-MacBook-Pro.local

Single space (no linebreak)

Comment





### **Exercise 1a**

- Reminder about exercises:
  - -Groups of three
  - -One person shares screen
  - -Solve cooperatively
  - -Screen-sharer switches for next exercise
  - -I will visit each group
- You have been given Linux cheat sheets





#### **Exercise 1a**

#### **Objectives:**

- You understand the basics of SSH
- You have a working cluster login configuration

#### <u>Tasks:</u>

- Log into the cluster with password
- Familiarize yourself with Linux console if necessary
- Set up an SSH config on your local PC
- Set up password-less login (key pair)
  - -Windows users: may skip adding

#### Note the following page!





### Exercise 1a

- Let's assign training users now
- You may use your own account instead
- Cluster address: <removed>

Login info Feb, 15/16: User: schulungXY PW: <removed>

(where XY is a number between 01-12, will be assigned during course)

- If bored, get creative
  - Try launching different programs
  - -Figure out how to get to the other login node

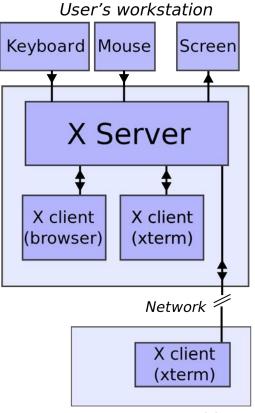
-...





### Linux Graphical User Interface

- X window system
- Basis of all Linux displays
- Can display windows from other computers
- X server needs to run on PC
- X <u>client</u> is software that window belongs to
- X windows can be transmitted by SSH connections



#### Remote machine





# Graphics via SSH

- Requirements
  - -X server installed on PC
  - SSH connection with X support
  - (Cluster supports X windows)
- Linux: X server built in
- Mac OS: Xquartz
- Windows: xming, MobaXTerm





#### Connecting with X support

- Enable X support in SSH
  - ssh -X <user>@<host>
  - -Must be upper case X
  - -Sometimes -Y used
    - "Trusted" connection
    - Less safe, sometimes necessary for things to work
- In config file: ForwardX11 yes or ForwardX11Trusted yes

Demo 5





# File Transfer

- Copying files between PC and cluster:
  - -Use scp command (secure copy)
- Syntax similar to Linux  ${\tt cp}$  command
- Uses SSH, can use same settings/presets
- Console-based, graphical front-ends also exist for all OSes





### File Transfer

#### • Syntax:

scp [options] sourcehost:sourcefile targethost:targetfile

- -Host may be left out if local
- -Host may be SSH preset
- -Source or target or both can be remote
- Same rule as cp about -r when copying entire directories
- Unlike cp: will print status of file transfer to screen
- Not only possibility (rsync)

Demo 6





### Third-party tools: connection

- Connecting: graphical clients exist for all OSes
- Windows: particularly important because native SSH support limited
  - Two main options:
    - MobaXTerm: modern, many features

-Integrated file transfer, X server, text editor, key generator

• PuTTY: trusted, only SSH connections (no X server)

-Separate X server: xming

• Mac OS: external X server necessary





# Third-party tools: file transfer

- File transfer: clients exist for all OSes
  - -Windows:
    - MobaXTerm
    - WinSCP
  - -Mac OS
    - Forklift
    - Cyberduck (no experience)

Demo 7





# Third-party tools

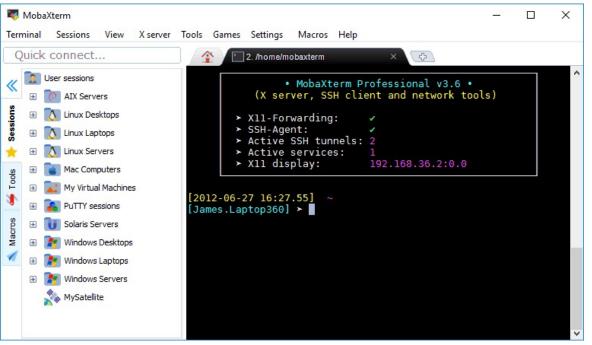
- Key point:
  - -All built on top of SSH and SCP
  - -Same concepts still apply
  - -Enter same login data
- MobaXTerm is particularly important
  - -Let's look at it more closely





#### Windows SSH Software

- MobaXTerm
  - -Free software (<u>mobatek.net</u>)
  - -All-in-one client
  - Does not need to be installed
  - -Specify host and user
  - -Good for newbies



Source: mobatek.net





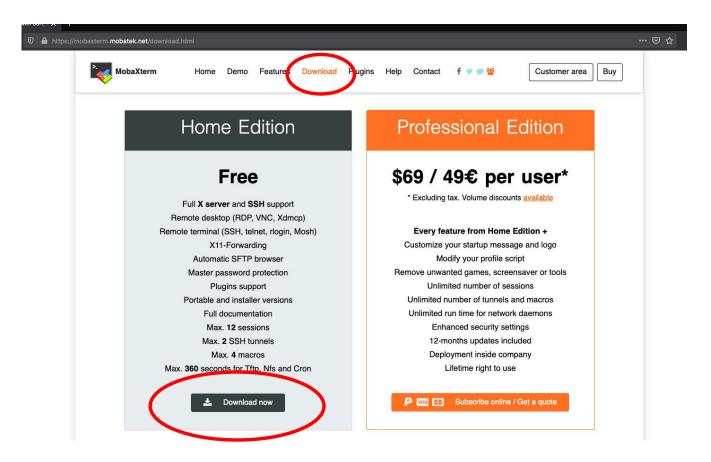
### MobaXTerm: Download

- Download MobaXTerm from <a href="https://mobaxterm.mobatek.net/">https://mobaxterm.mobatek.net/</a>
- Free
- Comes in "Installer" and "Portable" versions
  - -CIP Pools: download portable version, unzip, run .exe
  - -Cancel Windows firewall warning, it works anyway
- Windows users will do this in the first exercise





#### MobaXTerm: Download







#### Connecting: MobaXTerm

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	MobaXterm	Home	Demo	Features	Download	Plugins	Help	Contact	f 🦉 🗩 😫		Customer are	Buy	
	MobaXterm Home E	dition											
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	Download previous s	stable version	i: <u>Moba)</u>	Kterm Porta	ble v20.1 N	obaXterm	nstaller	<u>/20.1</u>					
	You can also get ear	ly access to t	the latest f	features and	-	its by down			Preview version	on:			
	By downloading Mot	aXterm softw	vare, you a	accept Mot	aXterm terms	and condi	ions						
	You can download M	lobaXterm an	nd plugins	sources he	re								
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#### Connecting: MobaXTerm

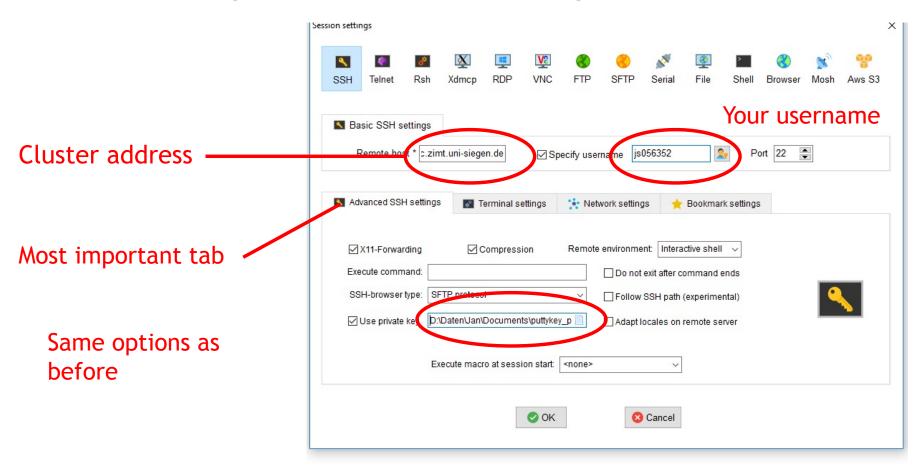
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February 15/16, 2022





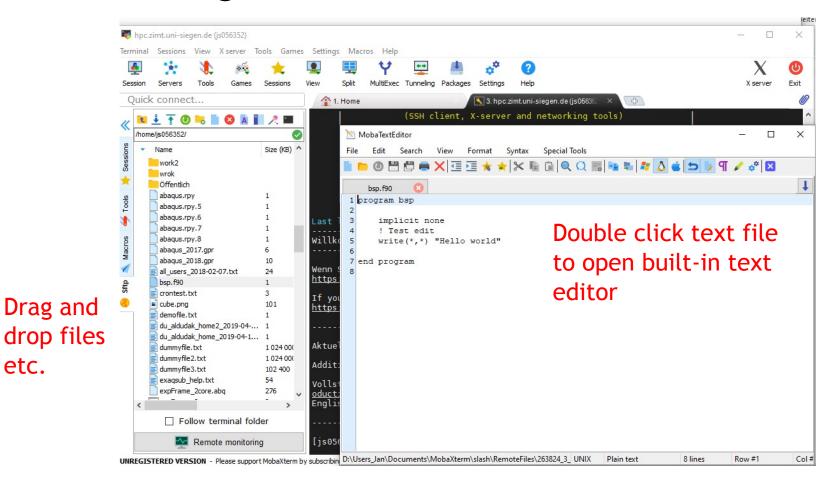
#### **Connecting: MobaXTerm configuration**







#### **Connecting: MobaXTerm features**



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etc.





# Outline

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  - Exercise 3: SLURM options





### **Exercise 1b**

#### **Objectives:**

- You are familiar with X servers and file transfer
- You have a setup you are comfortable with

#### <u>Tasks:</u>

- Set up SSH connection with X server
- Practice using SCP
- If on Windows: install MobaXTerm and set it up
- If on Mac OS: install Xquartz
- Play around with GUI clients of your choice

Remember: if bored, get creative





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### Using the cluster

- Key differences to regular PC
  - -Home vs. Workspaces
  - -Environment modules
  - Parallel programs/libraries
  - -Jobs





#### Workspaces

- /home usually limited in size (100 GB in our cluster)
- Workspaces for CFD data
  - -Higher bandwidth
  - -Unlimited storage (but limited in time)
  - -OMNI: /work
  - -Burst buffer: /fast
- Workspace mechanism: allocate for X days
  - -ws allocate <name> <days>
  - -ws\_list
  - -ws release <name>





### **Burst buffer**

- OMNI has so-called "burst buffer"
  - -Made up of SSDs
  - -Considerably faster file input/output
  - -32 TB
- Only use it if you need it
   Limited space for all users
- Works identically to normal workspaces
  - -Additional option ws\_allocate -F fast <name> <days>
  - -File system: /fast





#### Workspaces

- Additional options:
  - -Send e-mail before workspace expires
  - -Generate calendar item : ws\_send\_ical
- Maximum duration: 30 days
  - -ws\_extend <ws-name> <days>
  - -Can be extended up to 3 times
  - -Extensions and remaining time with ws\_list
- After that, data is GONE!
  - -Can be rescued by admins for 10 days after that
  - -Do not rely on this

Demo 8





#### Workspaces

- Common problems:
  - -Forgetting duration in ws\_allocate <name> (will result in 1 day duration)
  - -Forgetting to renew WS

- Tip: set up your e-mail address
  - -Put a file named .ws\_user.conf in your home directory
  - -Inside file: mail: <Your e-mail address>
    - Note space after colon (YAML syntax)
  - -When creating workspace: ws\_allocate -r <days>
  - You will get an e-mail <days> before expiration





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Excursion: what happens in Linux if you type a command?

- Linux looks for program with that name
- Directories where Linux looks: defined by PATH environment variable
  - -Directories set by Linux
  - -Directories added by installed software (so it gets found)
  - You can add your own
- Goes through in order listed in PATH
  - -First hit gets executed





- PATH is called an environment variable
- Other variables set by Linux, e.g.: HOME, USER
- Set by programs to find libraries etc.
- Used by SLURM
  - -Special variables inside job
  - -Used to provide job information
- "Environment" because process sees it, provides it to subprocesses





- Many users with different needs
  - -Different versions of same software/library
  - -Different software with same commands
- Reconfigure environment for every user?
- Better: modular environment

   Users load module that they need
- Example:

```
module load openmpi4
module avail
```





- Modules may be loaded as dependency
- Some modules are loaded on login for each cluster user
- module list shows loaded modules
- module purge unloads everything (e.g. debugging)
- Possible to define own modules (see website)

Demo 9





## Environment setup on OMNI

- OMNI cluster is multipurpose
- Software from several different sources
- Tricky module setup
- Default modules always loaded: SLURM, GCC compiler, OpenMPI –Some modules depend on specific compiler or MPI
- Modules come in four groups





## **Environment setup on OMNI**

- Software sources:
  - -CentOS (operating system)
  - -Bright (cluster management software)
  - -OpenHPC (software collection)
- OpenHPC modules are only displayed if compiler and MPI is correct
  - -Gnu GCC vs. Intel Compiler
  - -OpenMPI vs. IntelMPI
- Some modules do not depend on any of those
  - -GPU modules are even mutually exclusive

Demo 10





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## Running computations: jobs

- A single HPC computation is called a job
- Job Scheduler SLURM
  - -Manages when to run jobs
  - -Efficient usage of resources
  - –Several commands (each with -h for options)
- One job = one command/script
  - -Start job: srun (--pty) <options> <linux-command> sbatch <options> <scriptname>
  - -Monitor jobs: squeue, show partitions: sinfo
  - -Delete job: scancel <job-id>





#### Running computations: queues

- Jobs are put into queues (called partitions in SLURM)
  - Different runtime
  - -Different size
  - -Different type of node (e.g. GPU)
- Each queue has default values
- You pick queue, runtime, number of nodes

 $\rightarrow$  As many resources as necessary, as few as possible (with safety margin)





#### Running computations: queues

#### Primary queues:

- debug:
  - -Only for testing
  - -15 minutes runtime
- short, medium long
- expert:
  - -For big jobs
  - -Users must contact us and obtain permission





#### Running computations: queues

Special queues:

- gpu:
  - If you want to use GPUs
  - -Needs additional options in job script: --gres=gpu:X (where X is
    number of GPUs needed)
- smp:

-two nodes, 1536 GB RAM, 64 cores (Intel CPUs)

• htc:

- HPE Moonshot system

Demo 11





## Monitoring jobs

- Check regularly what your job does
  - Your first job will fail (guaranteed)
  - -Might be a bug later on
  - -Might be a problem with the cluster
  - -Might run out of resources
  - -Might not be finished when time limit is reached
- Main command to check what your job is doing: squeue
- If possible, use checkpointing (write intermediate results)





### Monitoring: squeue example

[js056352@login2 ~]\$ squeue		
JOBID PARTITION NAME USER ST TIME NOT	<b>DES</b>	NODELIST(REASON)
54791 defr Vort3-s gk637 PD 0:00	1	(Priority)
54404 Zong Saoa2.tx c.352 PD 0:00	12	(Resources)
Unique ID of job 54393 defg RCT4-con 3033441 PD 0:00	9	(Priority)
54739 short rry ak339 PD 0.00	2	(Resources)
mediu Status: 75 R 3:23	1	cn079
Joh script name 10 Mumber of podes	1	cn088
54493 Lor PD: Pending	1	cn097
54620 det R: Running 05 R 9:00:35	2	cn[045-046]
54721 mediur CD: Completed 84 R 1:56:21	2	cn[154-155]
54748 det 39 R 1.12.09	1	cn042
54743 det F: Fail 39 R 1:19:02	1	044
54796 det 40 R 3:03	1	cn080
53880 long LCURVO g+657 R 1-14:36:41	1	cn096
53879 long LCURV gf657 R 1-14:36:55	1	cn095
FARCE land IT Fr		cn102
52806 Long TS5sh		cn087
54757 short MA gk339 R 59:28		cn[017-018]
54756 short MA gk339 R 1:06:59		cn[038-039]





## Other SLURM commands

- srun can also be used within job
  - -Runs command once in every task
  - -Warning: scripts need to be executable
- squeue -u <Your Username> will list all your jobs
- sinfo will list available partitions, spartition lists defaults
- scancel <Job ID> will kill a job
   scancel -u <Your Username> kills all your jobs
- scontrol allows more in-depth information
   —Example: scontrol show job <Job ID>





## Other key concepts of SLURM

- SLURM allows you to choose how many and which resources to use
  - -Nodes
  - $-\mathsf{RAM}$
  - -Running time

• For now: one task = one program, using one CPU core

Demo 12





## Workflow: queuing a job script

- 1. You write the job script
  - Calls your software
  - Provides job settings
  - Loads environment
  - Any other necessary tasks
- 1. You prepare your software and files, workspace etc.
- 3. You queue your script with sbatch
- 4. You wait for job to complete, check intermediate results





#### Example job script

#### #!/bin/bash

- **#SBATCH** --time=0:20:00
- **#SBATCH** --nodes=1
- **#SBATCH** --tasks-per-node=6
- **#SBATCH** --mem 48000
- **#SBATCH** --partition=short

module load abaqus/2017

```
echo "Number of tasks: "
```

echo \$SLURM NTASKS

abq2017hf9 job=Test.inp mp mode=mpi interactive cpus=\$SLURM NTASKS





### Example job script

- #!/bin/bash
- **#SBATCH** --time=0:20:00
- **#SBATCH** --nodes=1
- **#SBATCH** --tasks-per-node=6
- **#SBATCH** --mem 48000

```
#SBATCH --partition=short
```

```
module load abaqus/2017
```

```
echo "Number of tasks: "
```

echo \$SLURM\_NTASKS

#### Which shell to use (Linux command)

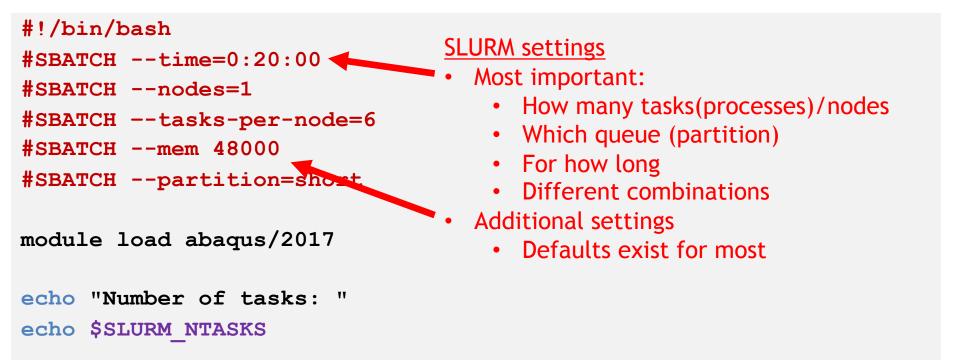
- At least two different families (csh,bash)
  - Different syntax
- Default on cluster: bash
- Does not have to be shell

abq2017hf9 job=Test.inp mp\_mode=mpi interactive cpus=\$SLURM\_NTASKS





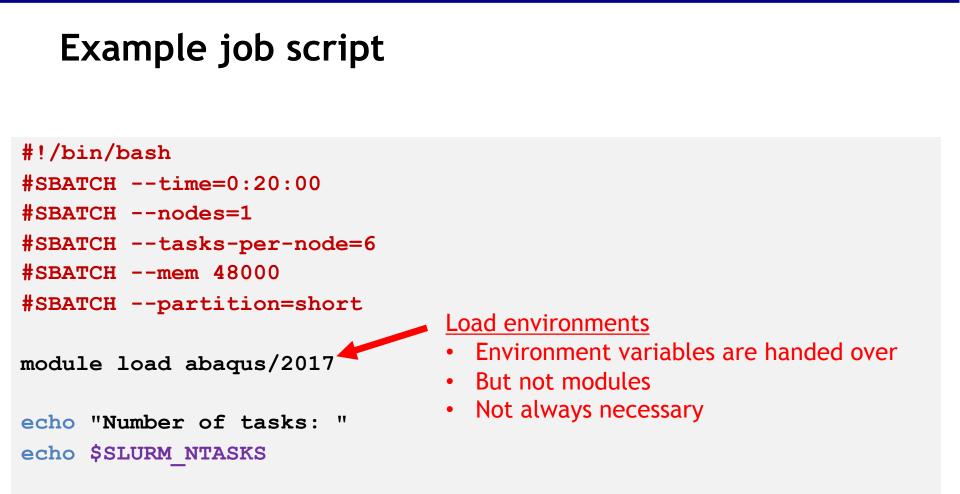
### Example job script



abq2017hf9 job=Test.inp mp\_mode=mpi interactive cpus=\$SLURM NTASKS







abq2017hf9 job=Test.inp mp\_mode=mpi interactive cpus=\$SLURM\_NTASKS



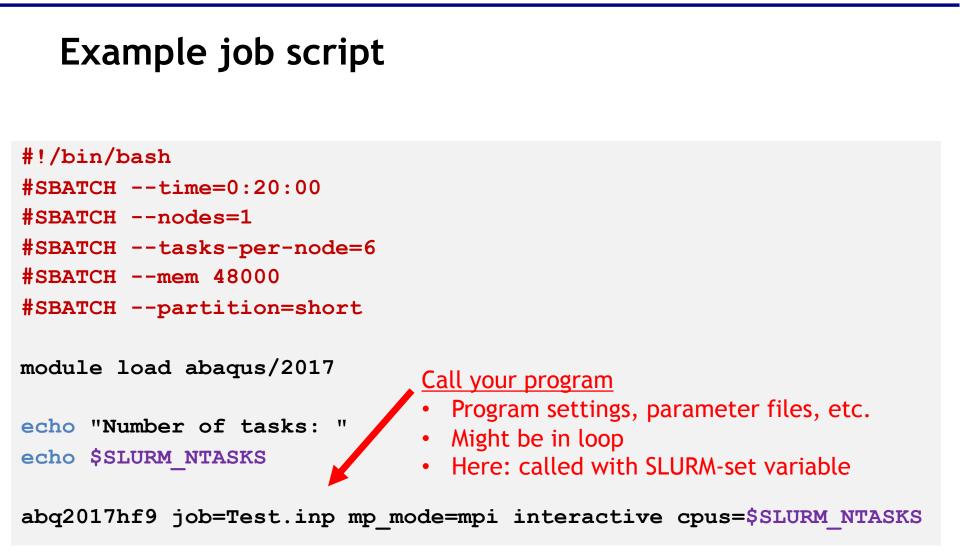




abq2017hf9 job=Test.inp mp\_mode=mpi interactive cpus=\$SLURM\_NTASKS



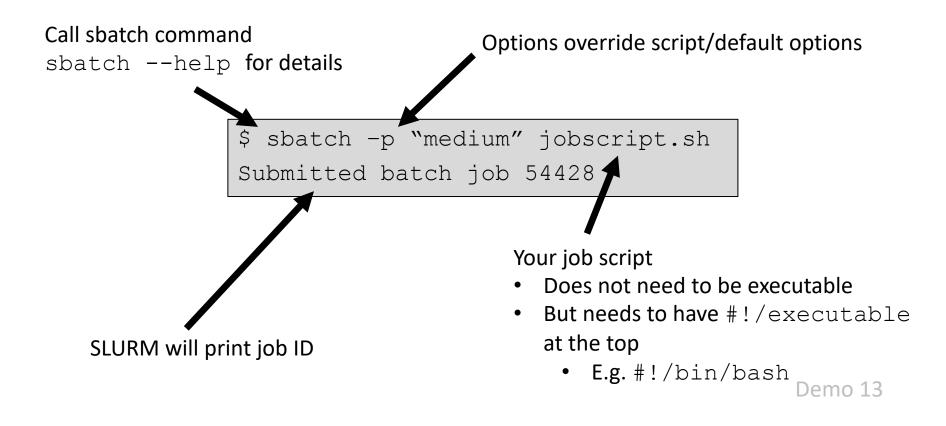








### Using sbatch to queue your job script







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## **Exercise 2**

#### **Objectives:**

- 1. You know how to create a simple job script
- 2. You can interpret the output of squeue and sinfo

#### <u>Tasks:</u>

- Write a job script that prints its working directory, sleeps for 30 seconds, then exits
  - -Remember the cheat sheets
  - You are allowed to google basic Linux commands

Note the following page!





## **Exercise 2**

- If bored, get creative:
  - -Use sinfo to find out how much of the cluster is currently busy
  - -Load and unload modules, use which command to see which program is called with a command
  - Try finding out file transfer speeds between your PC, your home directory and your workspace
  - -Try sbatch-ing a script in a different language

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- ...





### Solution: job script

```
#!/bin/bash
```

- **#SBATCH** --time=0:05:00
- **#SBATCH** --tasks=1
- **#SBATCH** --partition=short

```
# Print directory.
pwd
```

```
# Sleep.
```

```
sleep 30s
```





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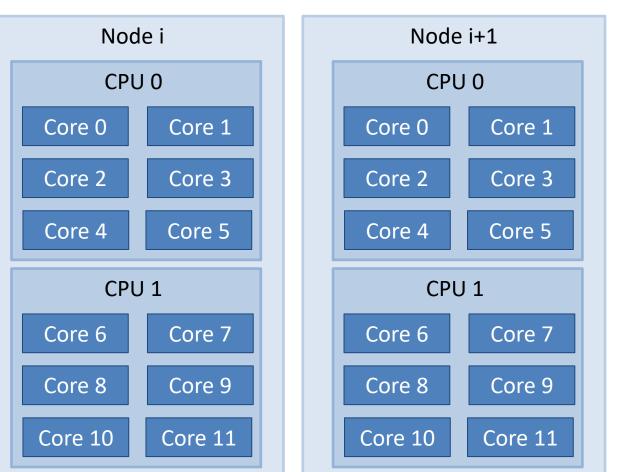




### Hardware visualized

Hardware:

- Cluster has <u>nodes</u>
- Nodes may have multiple <u>CPUs</u> (each on its <u>socket</u>), often 2
  - Not always important which CPU
- CPU has multiple cores





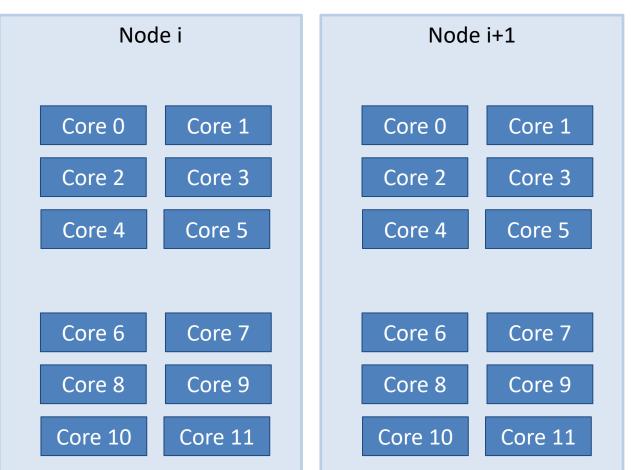


### Hardware visualized

#### **Simplification:**

- Difference between CPUs mostly matters for high-performance applications
- Communication between sockets is longer
- Separate caches

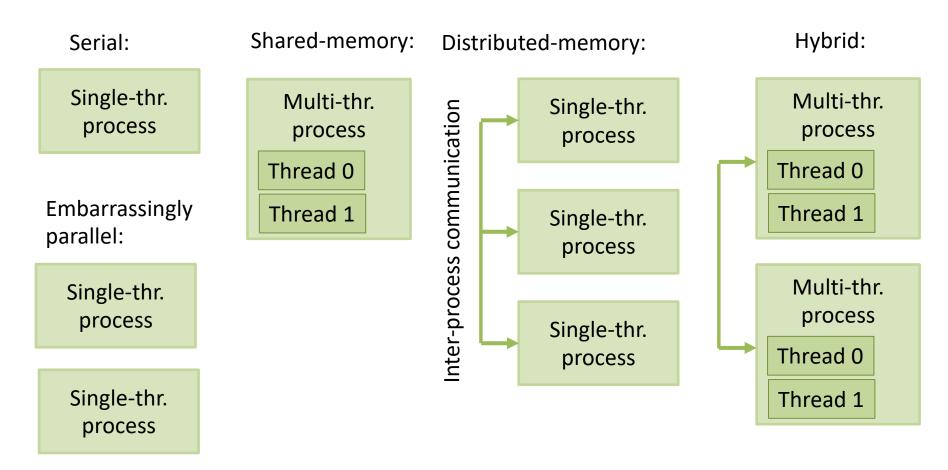
 $\rightarrow$  Ignored for now







## Workloads common in HPC





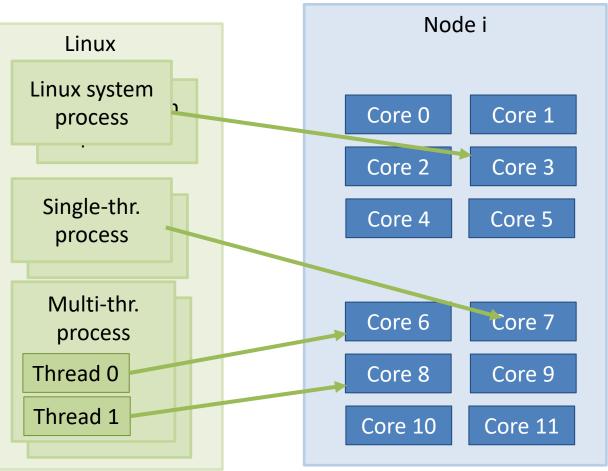


### Software visualized

#### **Operating system:**

- Each node is a separate computer
- OS runs processes
- Processes may have one or multiple threads

OS decides which process runs on which core(s)







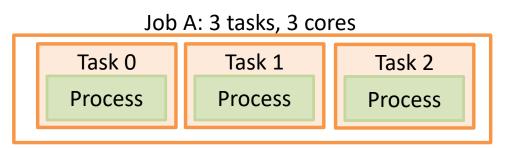
## Example SLURM jobs

 User specifies which and how many processes and threads to run in job

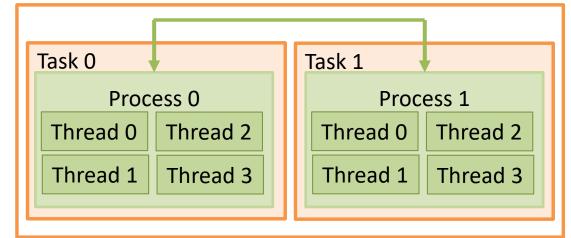
→ Tasks

#### What SLURM does:

- Decides on which nodes to run job
- Decides which nodes and processes a job gets
- Distributes tasks











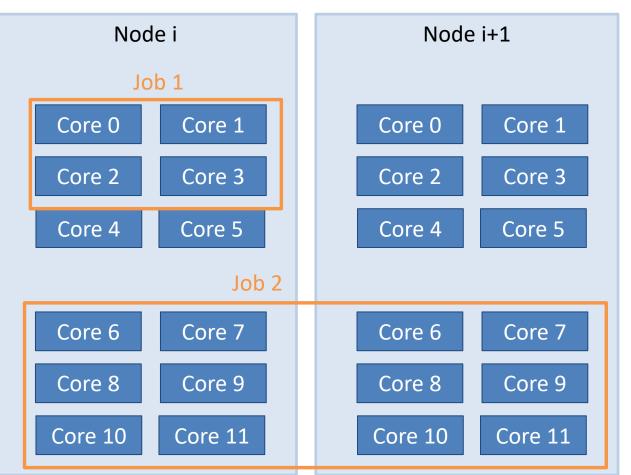
## Example SLURM jobs

 User specifies which and how many processes and threads to run in job

→ Tasks

#### What SLURM does:

- Decides on which nodes, cores to run job
- <u>Allocates</u> (reserves) resources
- Launches tasks



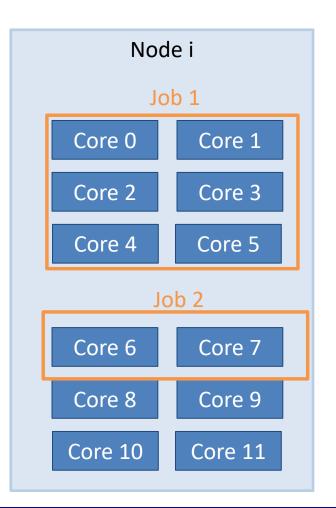




## Example SLURM jobs

#### **Complications:**

- Longer queues allow two jobs to share a node
  - Unusual for HPC clusters
- Annoying: SLURM refers to "cores" as "CPUs"
  - Again: more important for high performance applications
  - Don't get confused
  - Term "sockets" used when this is important







### What does this mean for you?

- Mostly, you need to decide how many tasks, how many CPUs per task
- Job options:

	<b>#SBATCH</b> ntasks=128	#SBATCH -n 128	
• Or alternatively:			
#SB	ATCHnodes=2	#SBATCH -N 2	2
#SE	ATCHntasks-per-nod	le=64	

• In both cases possible:

**#SBATCH** --cpus-per-task=4

Caution: sometimes starts with n, sometimes doesn't

• Consult sbatch documentation

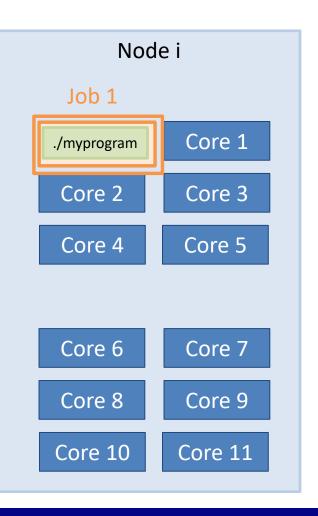




#### Simple serial program

- #!/bin/bash
- **#SBATCH** --time=0:20:00
- **#SBATCH** --ntasks=1
- **#SBATCH** --partition=short

./myprogram

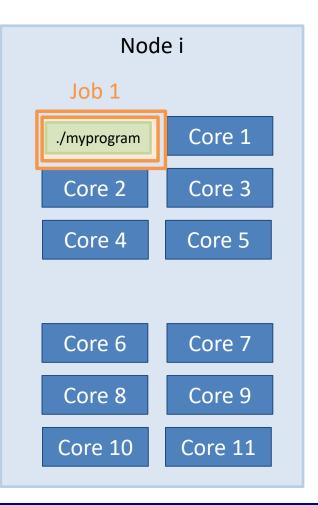






#### Simple serial program

- Simplest possible case
- Number of tasks = number of processes
- Default: only part of RAM used -RAM allocation covered later





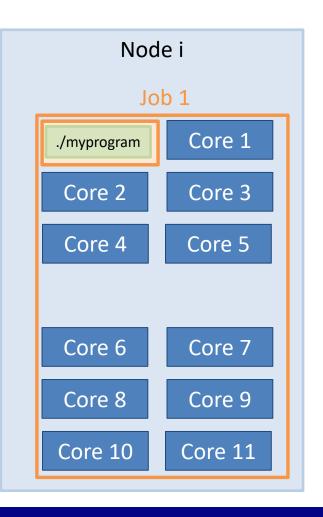


Wrong: serial program

#!/bin/bash

- **#SBATCH** --time=0:20:00
- **#SBATCH** --nodes=1
- **#SBATCH** --partition=short

./myprogram



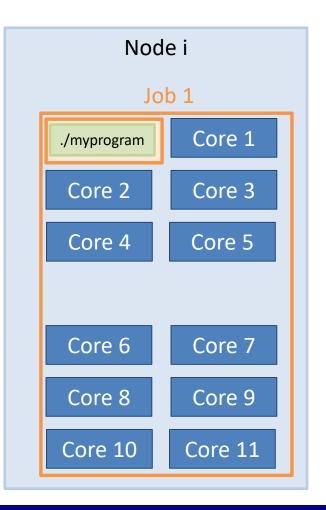




Wrong: serial program

• Waste of resources: full node allocated, only 1 task launched

- Might be legitimate reasons not to use all cores
  - -RAM allocation covered later
- But: allocation should reflect reality



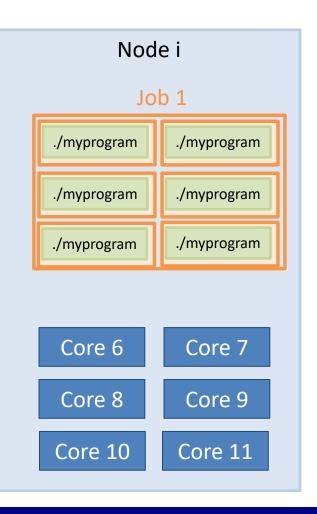




Many instances of serial program

- #!/bin/bash
- **#SBATCH** --time=0:20:00
- **#SBATCH** --ntasks=1
- **#SBATCH** --partition=short

srun ./myprogram

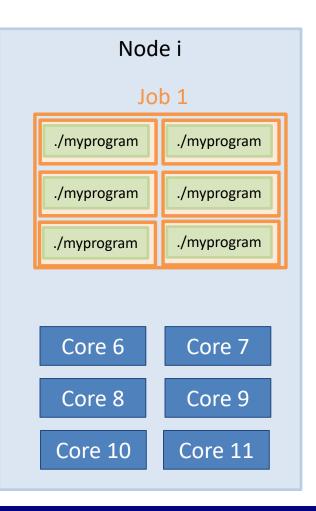






Many instances of serial program

- SLURM simply launches program multiple times
- Cannot talk to each other
- Be careful that they write to different files



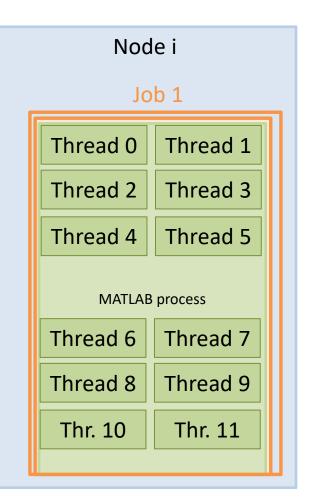




#### Launching MATLAB

- #!/bin/bash
- **#SBATCH** --time=0:20:00
- **#SBATCH** --ntasks=1
- **#SBATCH** --cpus-per-task=12
- **#SBATCH** --partition=short

matlab -nodisplay -r myscript

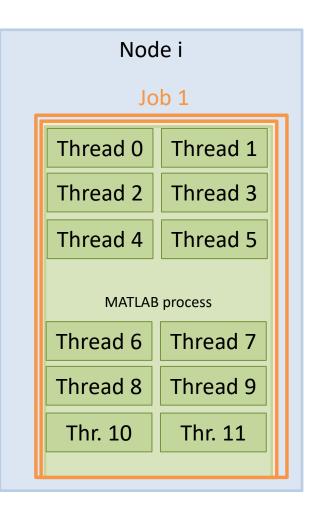






#### Launching MATLAB

- Commercial applications often multithreaded
- MATLAB: you do not even need to program differently
- Often good at auto-detecting which resources they have been given
- Does not use MATLAB advanced parallel features (pools), not covered







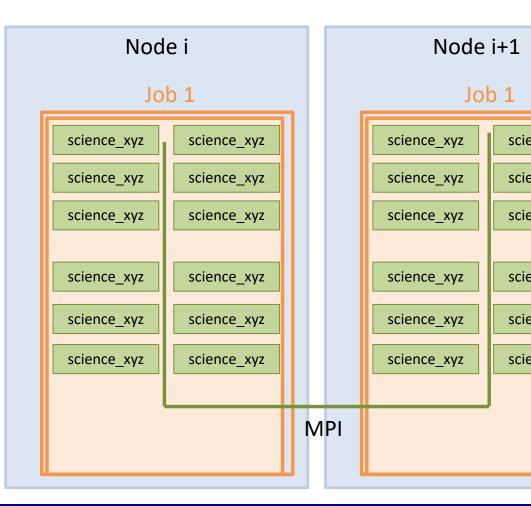
#### Launching scientific application

#!/bin/bash

- **#SBATCH** --time=4:00:00
- **#SBATCH** --nodes=20

**#SBATCH** --ntasks-per-node=12

mpirun -np 240 science\_xyz
parafile >log.txt

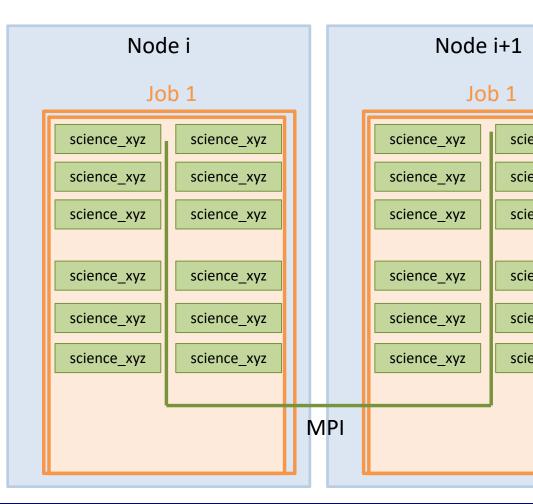






#### Launching scientific application

- Often lots of nodes
- Often distrib.-mem.
  - -Especially MPI
- Launch with mpirun -np [N]
- SLURM and MPI can generally talk to each other
- Documentation: "can also use srun instead of mpirun"
- Commercial: sometimes use internal MPI







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  - Jobs
  - Exercise 2: your first job script
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  - Tasks, processes, cores
  - Miscellaneous SLURM stuff
  - Exercise 3: SLURM options





## **SLURM defaults**

- SLURM has defaults for most options
  - Default queue (partition)
  - -Default runtime: spartition command to show
  - -Default task setup (caution)
    - 1 task, 1 CPU per task
- Find out what your job actually ran with:
  - -scontrol show job <Job ID>
  - -Only while running
- SLURM config readable to everyone: scontrol show conf

Demo 14





### SLURM: resources other than CPU cores

- So far we have only talked about CPUs
- SLURM does the same kind of management for RAM
- Default: ~4 GB RAM per task
  - -Reason: 64 cores, 256 GB RAM total (240 for apps)
  - -Option --mem 0 to allocate all RAM on node
  - -Alternative: --mem-per-cpu <amount in MB>
- Again, scontrol show job is your friend here





### More about the srun command

• srun command has multiple uses

- Inside a job:
  - -launch a process for every SLURM task

- Outside of job:
  - -launch a job running one Linux command (as opposed to a script)
  - -launch interactive job





## The srun command and interactive jobs

### <u>Scenario:</u>

- You want to use a CPU-intensive application yourself -e.g. visualization, post-processing
- Will slow down/block entire login node
   We reserve the right to kill processes
- You cannot start a batch job because application needs your input

### Solution:

- Interactive job
  - Resources allocated like any other job
  - -But only a console is opened and you can work within it





## The srun command and interactive jobs

#### Interactive job:

- 1. Use srun, not sbatch
- 2. Use --pty option
- 3. Use other SLURM options as needed
- 4. Specify which command (typically /bin/bash)
- 5. Wait for job to start (console stuck, then it opens)

srun --pty -t 5:00:00 /bin/bash





# **SLURM:** job priorities

- How does SLURM decide when your job runs?
   –Setup such that people do not have to wait too long
   –sprio command
- Priority for each job
  - -How long has it been waiting? (16%)
  - -How many core-hours has the user recently used? (80%)
  - -Bigger job are slightly preferred (4%)
- Additionally: "backfill" mechanism
  - -Plays "Tetris", fits small jobs onto free nodes

Demo 16





## Miscellaneous SLURM info

- Job arrays: multiple identical jobs

   Grouped, don't pollute queue
   Max 200 jobs per user on OMNI
- If you see "accounts" mentioned, not used on OMNI
- Remember environment variables inside job
- When in doubt: SLURM documentation is quite extensive -Many more options: run on specific nodes etc.





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## **Exercise 3**

### Objectives:

- 1. You understand the differences between SLURM task options
- 2. You can interpret scontrol show job output

### <u>Tasks:</u>

- Take your job script from earlier, try different combinations of parameters
  - -Also leave out parameters
  - Remember the cheat sheets + Google + SLURM documentation
- Check what your job actually did with squeue and scontrol show job

Note the following page!





## **Exercise 3**

- If bored, get creative:
  - -Find out how to queue a job with srun instead of sbatch
  - -Look into SLURM conf file with scontrol show conf. What do you recognize/not recognize?
  - -Google SLURM job parameters that you do not recognize
  - -Find out what happens if you try impossible parameters (e.g. 100 CPUs on one node)

-...





### Thank you for your attention

**Questions?** 

February 15/16, 2022

Introduction to the OMNI cluster