



## Reasons for using CSC computing resources



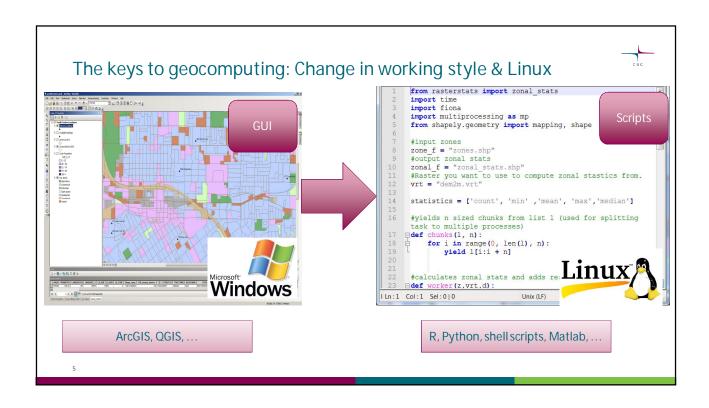
- Computing something takes more than 2-4 hours
- Need for more memory
- Very big datasets
- Keep your desktop computer for normal usage, do computation elsewhere
- Need for a server computer
- Need for a lot of computers with the same set-up (courses)
- Convenient to use preinstalled and maintained software
- Free for Finnish university and for state research insitute users

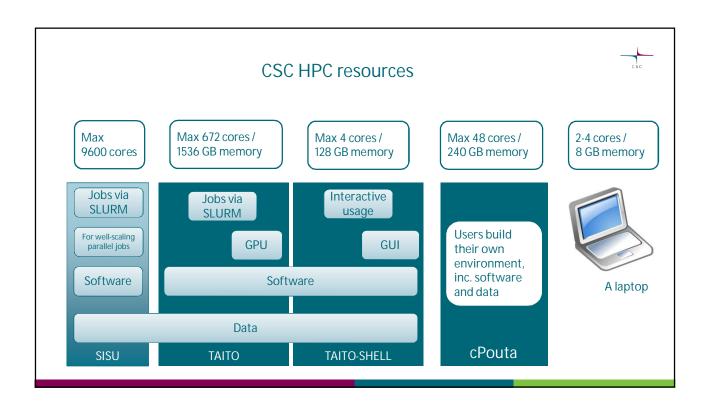
# CSC main computing resourcs for GIS users



- Taito supercluster
  - o Pre-installed software
  - o Some GIS data available
- Before using:
  - o Move your data and scripts
  - o Small extra adjustments for running your scripts

- cPouta cloud
  - o The user sets up the environment o More freedom, and more responsibilities
- Before using:
  - o Setting up the computer
  - o Installation of software
  - $\circ\, \text{Move your data and scripts}$





# Example GIS use cases Taito



- Satellite data analysis
- Forest modelling
- Prioritization of protected areas
- Climate variables modelling
- Spiecies modelling
- Routing / travelling times
- DEM analysis

Main limiting factor: software not suitable for Taito

# Example GIS use cases in cPouta



- Course environment
- Routing / travelling times
- Catchment area calculations on the fly (custom code, Leaflet)
- Satellite data analysis
- Drone image analysis with opendronemap
- Raster data management
- Sharing research results with GeoServer
- Data sharing GeoNode

#### Taito vs cPouta

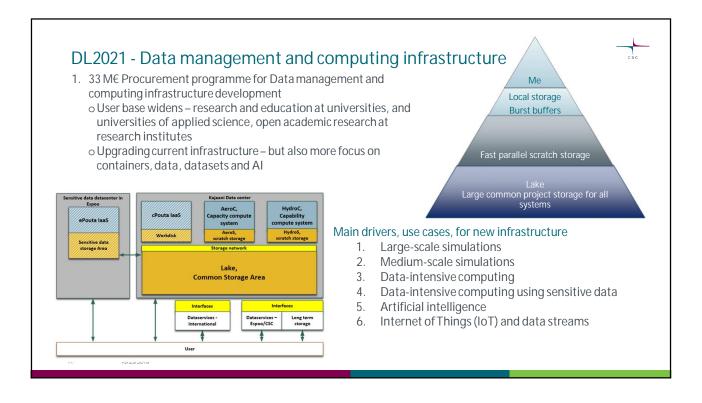


- Taito
  - o Ready working environment
  - o Preinstalled software
  - o Finnish GIS data
  - o Users created by CSC
  - o Limitations on software that can be used: Linux, no server, no root access
- cPouta
  - o Free hands, but also more responsibility
  - o Always starting from zero
    - $\circ \, \mathsf{Operating} \, \mathsf{system}, \, \mathsf{firewalls}, \mathsf{security} \, \mathsf{groups}, \, \mathsf{software} \, \mathsf{installation}, \mathsf{users} \, \mathsf{etc}$
    - o Back-up

# Realistic expectations



- A single core of a CSC machine is about as fast as one of a basic laptop.
- It has just a lot of them.
- .. and more memory and faster input-output.
  - > Just running your single core script at CSC does not make it much faster.
  - ➤ For clear speed-ups you have to use several cores.
  - ➤... or optimize your script.



### Main new features for users

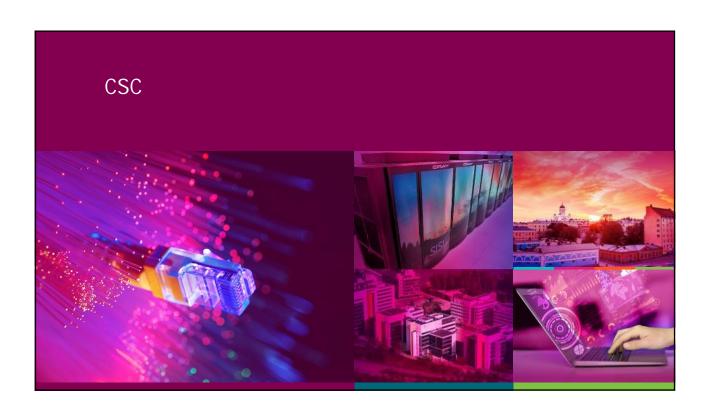


- More memory per node
- Big object storage area
- Running Docker-like containers in "Taito"
- More capacity -> shorter queues for batch jobs
- More modern system libraries -> less problems with software installation

# Timetable

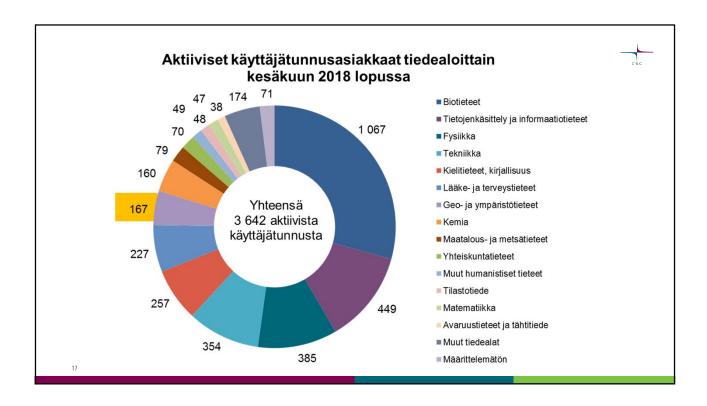


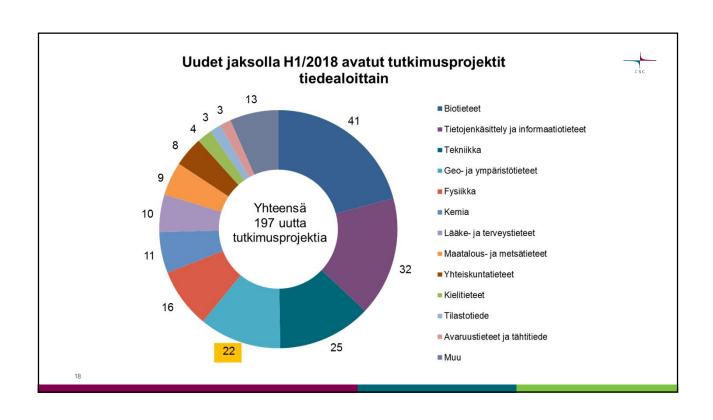
- The provider of the new system should be clear in autumn 2018
  New "Taito" in 1H/2019
- New "Sisu" in 2020

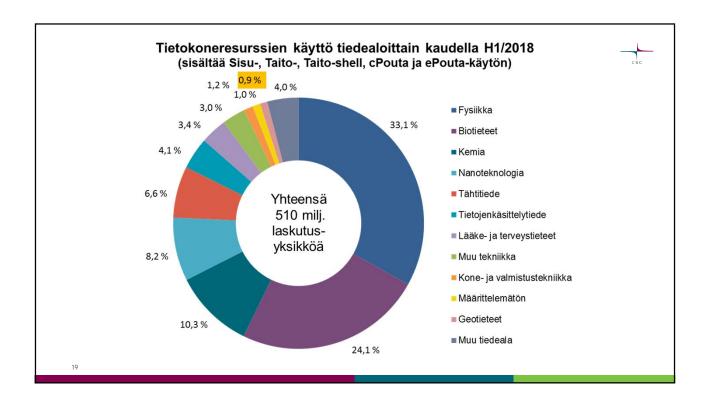










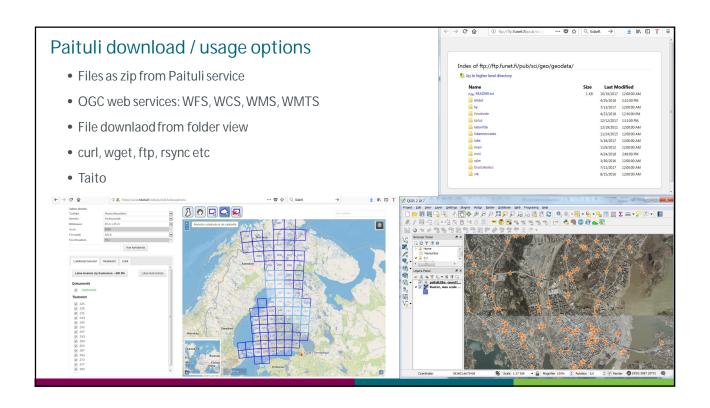




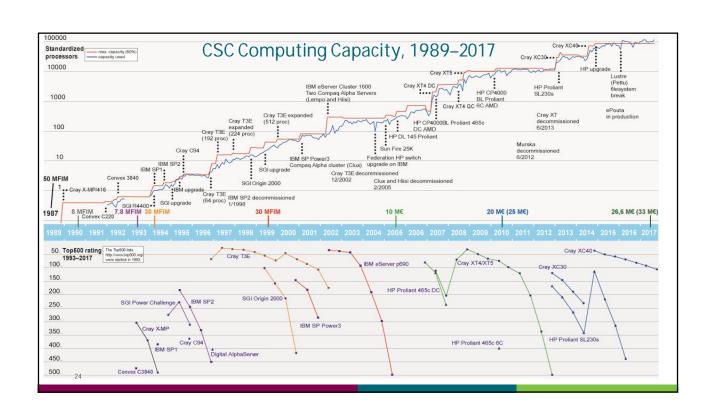
#### Paituli

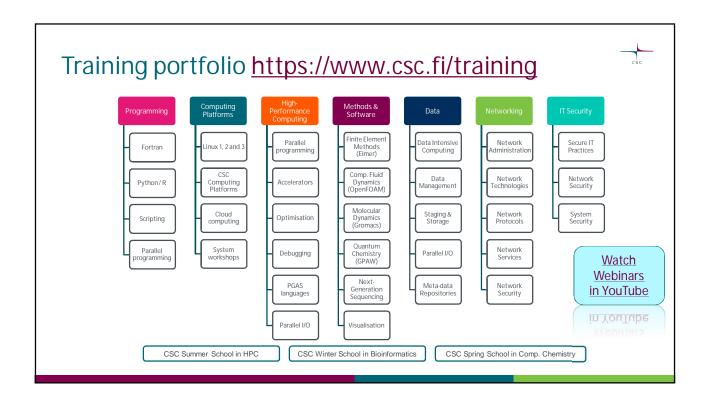
- www.csc.fi/paituli
- File download service of spatial data
- Datasets sharing service for scientific spatial data
- Almost all datasets are open to everybody
- In total 8+ Tb data.
- Includes historical versions of data (2005 ->)
- CSC operates the service, Ministry of Culture and Education supports the service, free of charge for users

# Paituli datasets producers Agency for Rural Affairs Finnish Meteorological Institute Finnish environmental administration Institute for the Languages of Finland Latuviitta National Land Survey of Finland Natural resources institute Finland Population register center Road Administration of Finland Statistics Finland University of Helsinki



# Four calls every year until April 2021 Visit a group or invite a (coming) collaborator to your group Next DL for applications is 15th November 2018 Requirements: Non-proprietary research Affiliated at an EU-country, associated country or other Project needs/benefits of HPC resources Provided: Support for accommodation, travel costs and likely a small daily allowance for a 3-13 week visit Resources and support from local HPC-center http://www.hpc-europa.org/



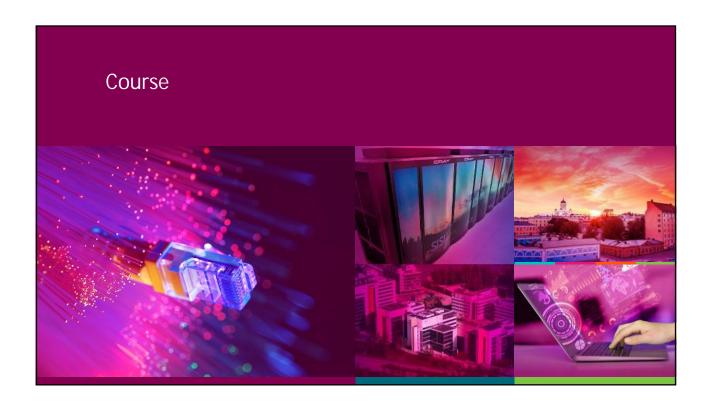


# GIS training 2018/19

# https://www.csc.fi/training



- 24-25.10.2018 Geocomputing with CSC computing resources
- 12-14.11 Introduction to GIS Python
- (15.11 Using Taito and R for spatial analysis, Turku)
- Google Earth Engine
- Geospatial data analysis with R
- Lidar data analysis
- Webinar: Paituli, cPouta for GIS
- Archive includes material of past courses



# Program, 24.10.2018



08:00 - 8:50 Linux intro

09:00–09:15 Introduction to the course

09:15–09:45 Getting access
User account, project and services,
web-based access to CSC's services

09:45-10:00 Coffee break

10:00–11:00 How to connect

How to access CSC's computers, NX
client, taito-shell

11:00–12:00 CSC's computing
environment Different platforms,
module system, licensing, storage and
data transfer

12:00-13:00 Lunch break

13:00–14:30 Running your jobs, resourcemanagement (a.k.a. batch job) systems

14:30-14:45 Coffee break

14:45–15:30 Running R code in Taito

### Program, 25.10.2018



09:00 - 10:30 Running Python code in Taito

10:30 - 10:45 Coffee break

10:45 - 12:00 GIS software and data in Taito. Using virtual rasters.

12:00 - 13:00 Lunch break

Taito

13:00 - 14:30 CSC's cloud services, and cPouta

14:30 - 14:45 Coffee break

14:45 - 16:00 Running GIS software in cPouta

cPouta

#### **Practicalities**



- Keep the name tag visible
- Lunch is served in the same building

   Room locked during lunch (lobby open, use lockers)
- Toilets are in the lobby
- Network:
  - o WIFI: eduroam, Haka authentication
  - o Ethernet cables on the tables
  - o CSC-Guest accounts
- Username and password for workstations: given on-site

- Bus stops
  - $\circ$  Other side of the street (102,103)  $\rightarrow$  Kamppi/Center
  - o Same side, towards the bridge (194,195/551) → Center/Pasila
  - o Bus stops to arrive at CSC at the same positions, just on opposite sides
- If you came by car: parking is being monitored - ask for a temporary parking permit from the reception (tell which workshop you're participating)
- Visiting outside: doors by the reception desks are open

#### Teachers on the course

#### Kylli Ek

- MSc in Geography
- Support / organizing
- Data management
- Softare development
- Paituli, ArcGIS/MapInfo licenses
- Taito
- Tools in use: GeoServer, PostGIS, Tools in use: GeoServer, OpenLayers, GDAL/OGR, QGIS, Etsin, Github
- JavaScript, R, Python, shell scripts

#### Eduardo Gonzalez

- MSc in Forestry
- Support / organizing
- cPouta and Rahti development
- Training development
- Notebooks
- Taito
- PostGIS, OpenLayers, GDAL/OGR, QGIS, Github
- JavaScript, R, Python, shell scripts

#### Elias Annila

- BSc in Geography
- Taito
- Tools in use: GDAL/OGR, QGIS, Github
- JavaScript, R, Python, shell scripts

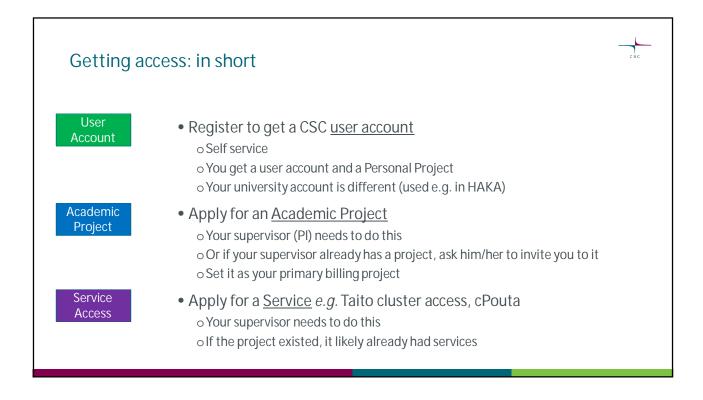
# **Participants**

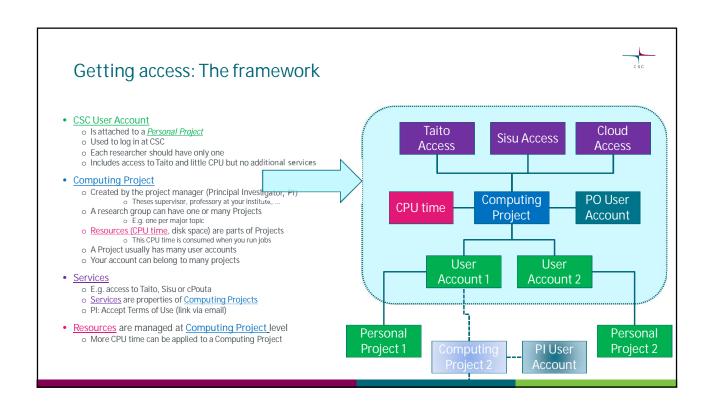


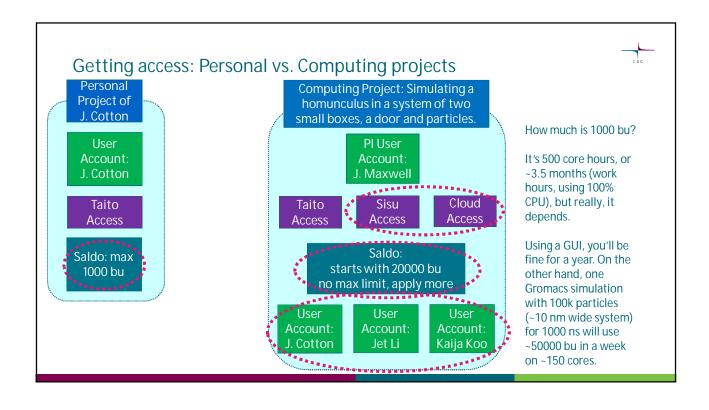
- Name
- Organization
- Interests in geocomputing
- Goals for the course

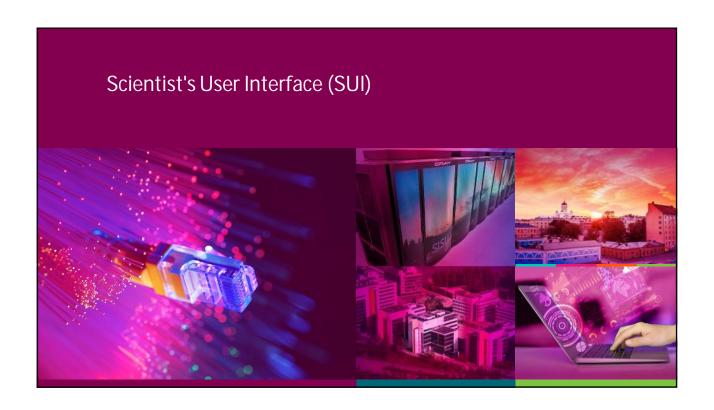


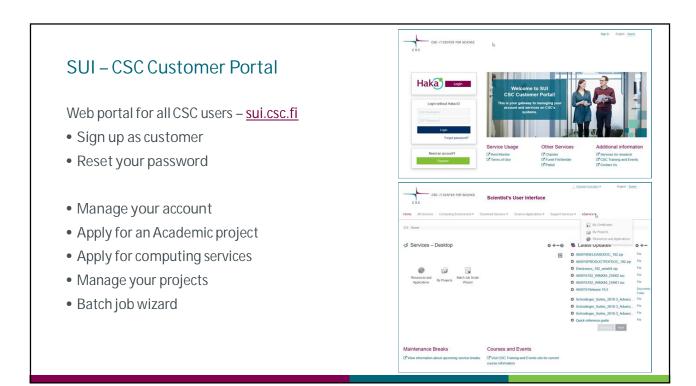


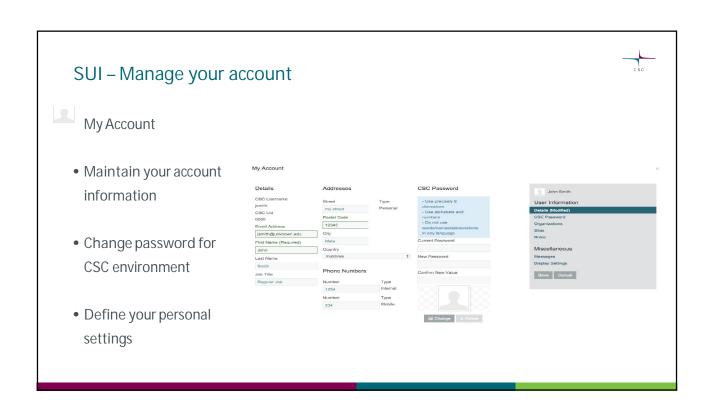


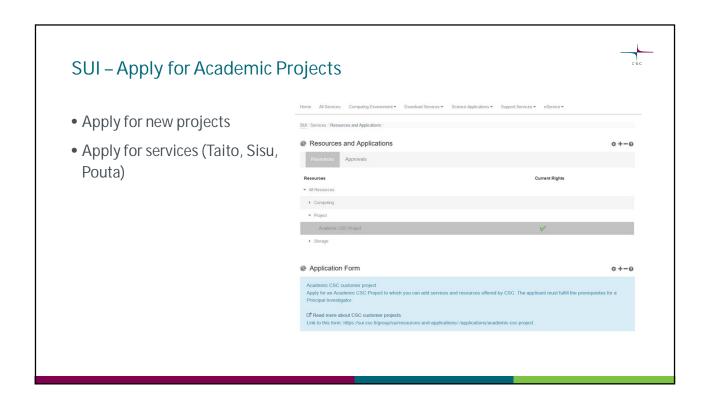


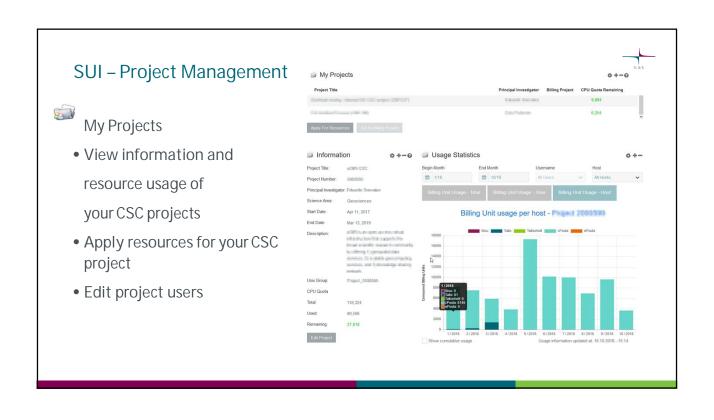


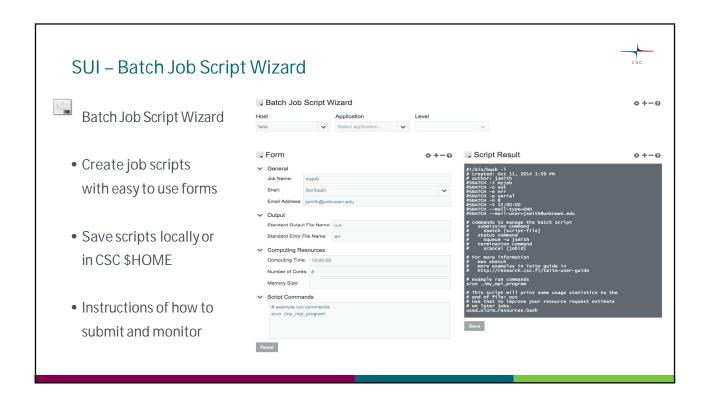














#### Direct ssh connection –Linux/Mac

CSC

- From UNIX/Linux/OSX command line
- Use –X (or –Y) to enable remote graphics\*
- scp: copy file to remote machine

# \$ ssh -X yourid@taito.csc.fi

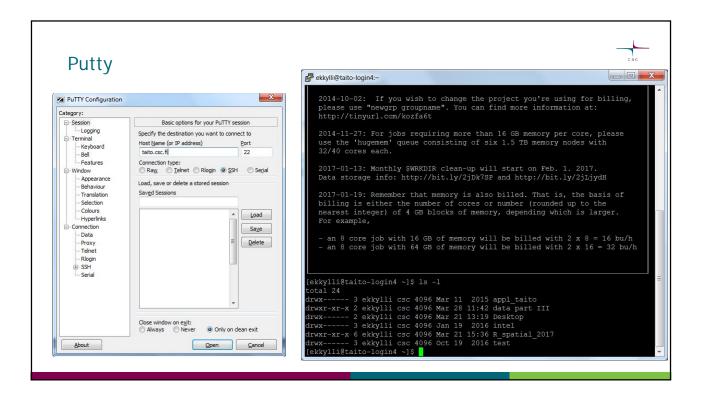
\* In Windows you'd also need an X-windows emulator, but there is a better way

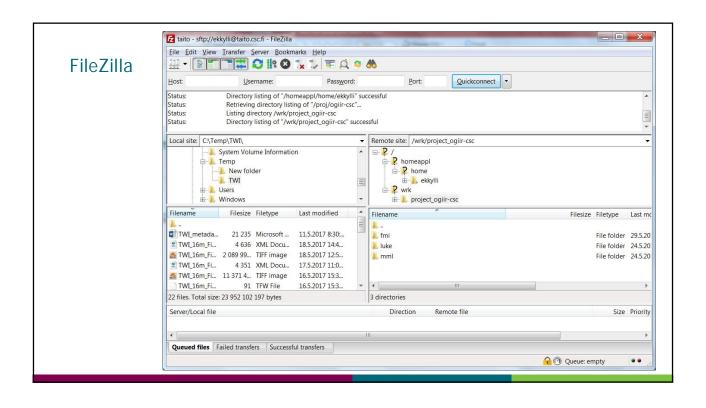
#### **Access from Windows**



- <u>Putty</u> for ssh connection
   o Can be installed without admin privileges
- NoMachine for GUI
  - o Needs Admin privileges for installation and update o Recommended method (also for Linux/Mac)
- FileZilla/WinSCP for moving data
   Efficient GUI
- Find about other access options and more information at: <a href="https://research.csc.fi/taito-connecting">https://research.csc.fi/taito-connecting</a>







# Tips for moving files



- When downloading from exernal services try to download directly to CSC, no via your local PC
- When lot of/big files use:
  - o rsync, if supported
  - o wget/curl, if only HTTP-urls
- Webinar Recording on Data Transfer

https://research.csc.fi/csc-guide-moving-databetween-csc-and-local-environment

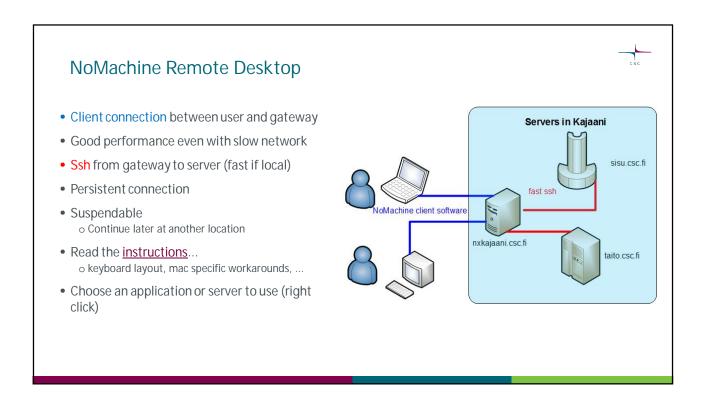
49

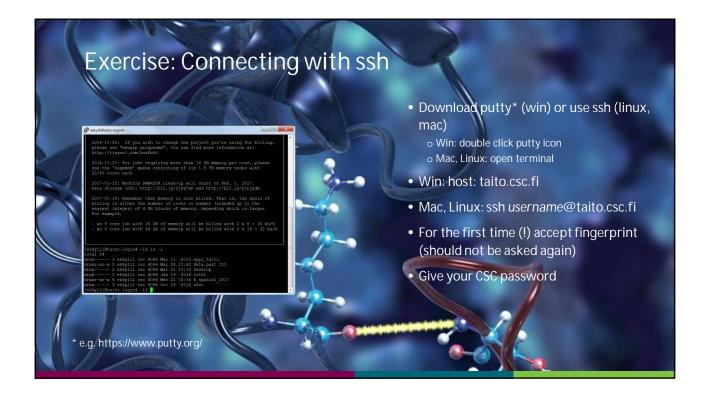
# Data transfer speed

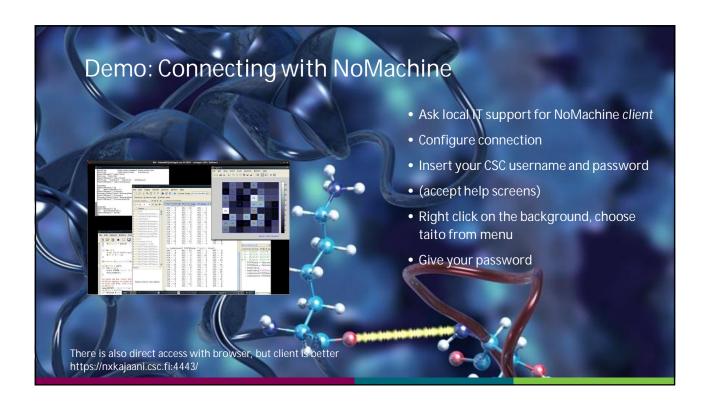


	50 Mb / s	25 Mb / s	5 Mb/s
1 kb	0,00002s	0,00001s	0,0002 s
1 Mb	0,02 s	0,01 s	0,2
1 Gb	20 s	40 s	3,5 min
1 Tb	5,5 h	11 h	2d 7h

- 50 Mb/s often the realistic fast speed
- 25 Mb/s good normal speed
- 5 Mb/s realistic speed in mobile network

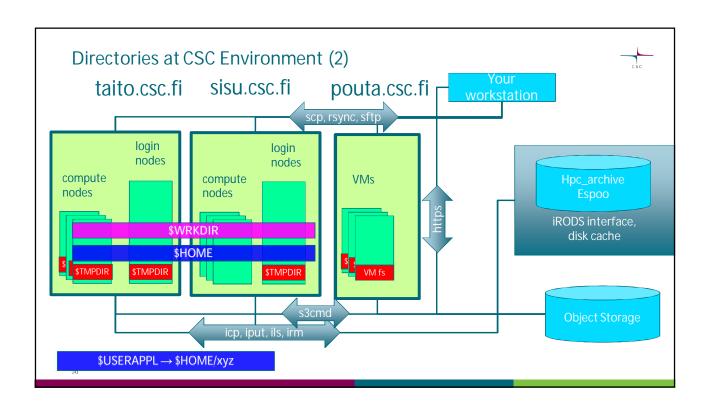








https://research.csc.fi/data-environment						
Directory or storage area	Intended use	Default quota/user	Storage time	Backup		
\$HOME <sup>1</sup>	Initialization scripts, source codes, small data files. Not for running programs or research data.	50 GB	Permanent	Yes		
\$USERAPPL 1	Users' own application software.	50 GB	Permanent	Yes		
\$WRKDIR <sup>1</sup>	Temporary data storage.	5 TB	90 days	No		
\$WRKDIR/DONOTREMOVE	Temporary data storage.	Incl. in above	Permanent	No		
\$TMPDIR <sup>3</sup>	Temporary users' files.	-	~2 days	No		
Project <sup>1</sup>	Common storage for project members. A project can consist of one or more user accounts.	On request	Permanent	No		
HPC Archive <sup>2</sup>	Long term storage.	2 TB	Permanent	Yes		
IDA <sup>2</sup>	Storage and sharing of stable data.	On request	Permanent	No, but multipl storage copies		





# The module system

csc

- Tool to set up your environment
  - o Load libraries, adjust path, set environment variables
  - o Needed on a server with hundreds of applications and several compilers etc.



# Typical module commands



module avail shows available modules (compatible modules in taito)

module spidershows all available modules in taitomodule listshows currently loaded modules

module load <name> loads module <name> (default version)
module load <name/version> loads module <name/version>

module switch <name1> <name2> unloads module name1 and loads

module name2

module purge unloads all loaded modules

Taito has "meta-modules" named e.g. r-env, which will load all necessary modules needed to run R.

# Module example



Simple plotting in R

> a = seq(0,10,by=0.1)

> plot(a,cos(a))

- Show compatible modules on Taito
  - \$ module avail
- Initialize R and RStudio statistics packages
  - \$ module load r-env
  - \$ module load rstudio
- Start RStudio using the command
  - s rstudio
- It's better to run the GUI (and calculations) on a compute node (jobs that have used 1h of CPU on the login node will be killed automatically)
- For interactive work, use taito-shell.csc.fi





# Batch jobs learning target

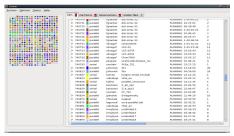


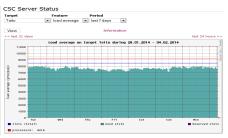
- Benefits of batch jobs for compute intensive jobs
   Difference of login and compute node
   Difference of interactive jobs (taito-shell) and batch jobs
- How to submit and monitor jobs
- Batch script contents *i.e.* resource requirements
- How to learn resource requirements of own jobs
- What is saldo [billing units]
- Submit first job(s)
- Learn to read the the manual

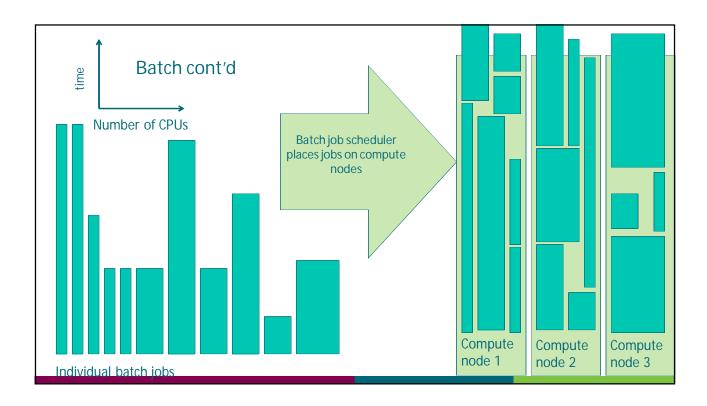
# What is a batch system?

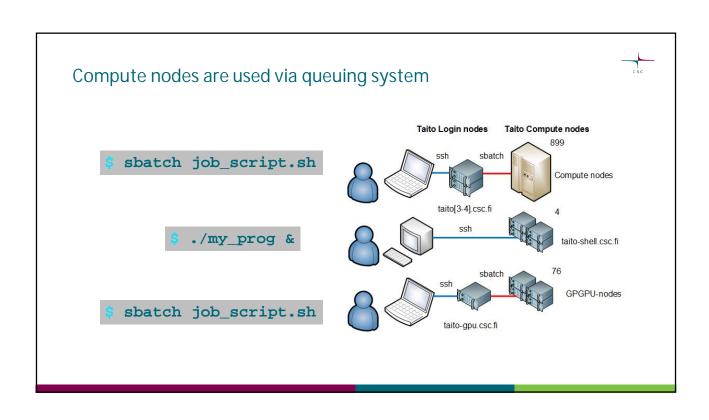


- Optimizes resource usage by filling the server with jobs
- Cores, memory, disk, length, ...
- Jobs to run are chosen based on their priority
- Priority increases with queuing time
- Priority decreases with recently used resources
- Short jobs with little memory and cores queue the least
- CSC uses SLURM (Simple Linux Utility for Resource Management)

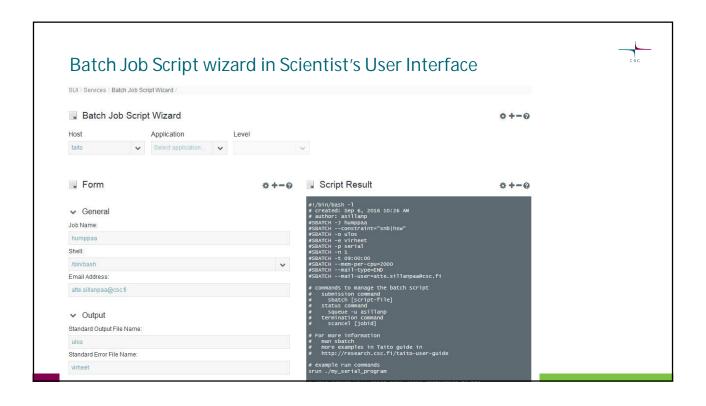












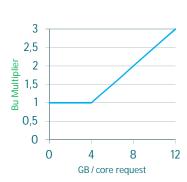
# Batch jobs: what and why



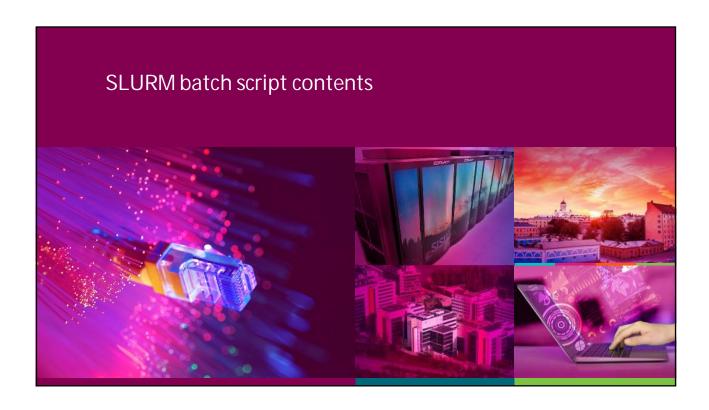
- > User has to specify necessary resources
  - Can be added to the batch job script or given as command line options for sbatch (or a combination of script and command line options)
- Resources need to be adequate for the job
  - Too small memory reservation will cause the job to fail
  - When the time reservation ends, the job will be terminated whether finished or not
- But: Requested resources can affect the time the job spends in the queue
  - Especially memory reservation (and perhaps requested time)
  - Using more cores does not always make the job run faster check!
  - Don't request extra "just in case" (time is less critical than memory wrt this)
- > So: Realistic resource requests give best results
  - Not always easy to know beforehand
  - Usually best to try with smaller tasks first and check the used resources
  - > You can check what was actually used with the seff command

# Saldo and billing units

- All jobs consume saldo
- https://research.csc.fi/saldo
- One core hour of computing equals 2 billing units [bu] times a multiplier
- Jobs requesting less than 4GB of memory per core have a multiplier of 1
- Jobs requesting 4GB or more per core have a multiplier X/4, where X is the requested memory per core:
  - o 5GB/core = 5/4x = 1.25x
  - o 12GB/core = 12/4x = 3x, ...
- Requested but not used computing time is not billed
- Requested but not used memory is billed
- If saldo runs out, no new jobs are possible
- New saldo can be requested from sui.csc.fi
- · GPU resources have an additional multiplier



- Serial job (1 core), 0.5 GB/core of memory, requested 24 hours, used 5 hours → billed: 1\*5\*2\*1=10 bu
- (failed) parallel job: requested 24 cores,
   2GB/memory per core, actually used 6 cores
   (18 cores idle) total run time 10 hours →
   billed 24\*10\*2\*1=480 bu
- Parallel job 3 cores, 5 GB/core, 10 hours → billed: 3\*10\*2\*5/4=75 bu



# Example serial batch job script on Taito



```
#!/bin/bash -l
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial
#SBATCH --constraint=snb

module load myprog
srun myprog -option1 -option2
```

#### #!/bin/bash -1

- Tells the computer this is a script that should be run using bash shell
- Everything starting with "#SBATCH" is passed on to the batch job system (Slurm)
- Everything (else) starting with "# " is considered a comment
- Everything else is executed as a command

```
pash -1
```

#### #!/bin/bash -1

```
#SBATCH -J myjob

#SBATCH -e myjob_err_%j

#SBATCH -o myjob_output_%j

#SBATCH --mail-type=END

#SBATCH --mail-user=a.user@foo.net

#SBATCH --mem-per-cpu=4000

#SBATCH -t 02:00:00

#SBATCH -n 1

#SBATCH -p serial
```

module load myprog
srun myprog -option1 -option2

#### #SBATCH -J myjob

- Sets the name of the job
- When listing jobs e.g. with squeue, only 8 first characters of job name are displayed.



```
#!/bin/bash -1
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial

module load myprog
srun myprog -option1 -option2
```

```
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
```

- Option -e sets the name of the file where possible error messages (stderr) are written
- Option -o sets the name of the file where the standard output (stdout) is written
- When running the program interactively these would be written to the command promt
- What gets written to stderr and stderr depends on the program. If you are unfamiliar with the program, it's always safest to capture both
- %j is replaced with the job id number in the actual file name

```
#!/bin/bash -1
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-
    user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial
module load myprog
srun myprog -option1 -option2
```

```
#SBATCH --mail-type=END
#SBATCH --mail-user=a.user@foo.net
```

- Option --mail-type=END = send email when the job finishes
- Option --mail-user = your email address.
- If these are selected you get a email message when the job is done. This message also has a resource usage summary that can help in setting batch script parameters in the future.
- To see actually used resources try also: sacct -1 -j <jobid> (more on this later)

```
#!/bin/bash -l
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
```

#SBATCH -o myjob\_output\_%j
#SBATCH --mail-type=END
#SBATCH --mail-user=a.user@foo.net

#SBATCH --mem-per-cpu=4000

#SBATCH -t 02:00:00

#SBATCH -n 1 #SBATCH -p serial

module load myprog
srun myprog -option1 -option2



#### #SBATCH --mem-per-cpu=4000

- The amount of memory reserved for the job in MB
  - 1000 MB = 1 GB
- Memory is reserved per-core basis even for shared memory (OpenMP) jobs
  - For those jobs it is better to ask memory per job:
  - --mem=1000
- Keep in mind the specifications for the nodes. Jobs with impossible requests are rejected (try squeue after submit)
- If you reserve too little memory the job will be killed (you will see a corresponding error in the output)
- If you reserve too much memory your job will spend much longer in queue and potentially waste resources (idle cores)

```
#!/bin/bash -1
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial
```

module load myprog
srun myprog -option1 -option2

#### #SBATCH -t 02:00:00

- Time reserved for the job in hh:mm:ss
- ➤ When the time runs out the job will be terminated!
- With longer reservations the job may queue longer
- Limit for normal serial jobs is 3d (72 h)
  - if you reserve longer time, choose "longrun" queue (limit 14d)
  - In the longrun queue you run at your own risk. If a batch job in that queue stops prematurely no compensation is given for lost cpu time
  - In longrun you likely queue for a longer time: shorter jobs and restarts are better (safer, more efficient)
- Default job length is 5 minutes → need to be set by yourself.

TIP: If you're unsure of the syntax, use Batch job wizard in <u>SUI</u>

#!/bin/bash -1

#SBATCH -J myjob
#SBATCH -e myjob\_err\_%j

#SBATCH -o myjob\_output\_%j

#SBATCH --mail-type=END

#SBATCH --mail-

user=a.user@foo.net

#SBATCH --mem-per-cpu=4000

#SBATCH -t 02:00:00

#SBATCH -n 1

#SBATCH -p serial

module load myprog

srun myprog -option1 -option2

#### #SBATCH -n 1

- Number of cores to use. More than one means parallel.
- It's also possible to control on how many nodes your job is distributed. Normally, this is not needed. By default use all cores in allocated nodes:
  - --ntasks-per-node=16 #(Sandy Bridge)
  - --ntasks-per-node=24 #(Haswell)
- Check documentation: <a href="http://research.csc.fi/software">http://research.csc.fi/software</a>
  - There's a lot of software that can only be run in serial
- OpenMP applications can only use cores in one node

```
#!/bin/bash -1
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-
   user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial
module load myprog
```

srun myprog -option1 -option2

#### #SBATCH -p serial

- The queue the job should be submitted to
- Queues are called "partitions" in SLURM
- You can check the available queues with command

#SBATCH -t 02:00:00 #SBATCH -n 1 #SBATCH -p serial module load myprog srun myprog -option1 -option2 STATE NODELIST draining c623 mixed c[25,76-77,. allocated c[3-24,26-75,... idle c[211-213,... draining c623 mixed c[25,76-77, idle c[211-213,... draining c623 mixed c[25,76-77,... idle c[211-213,.

#!/bin/bash -1 #SBATCH -J myjob #SBATCH -e myjob\_err\_%j

#SBATCH -o myjob\_output\_%j #SBATCH --mail-type=END #SBATCH --mail-user=a.user@foo.net #SBATCH --mem-per-cpu=4000

```
sinfo -1
[asillanp@taito-login4 ~]$ sinfo -l
Wed Jan 28 15:45:39 2015
PARTITION AVAIL TIMELIMIT JOB_SIZE ROOT
                                            SHARE
                                                      GROUPS NODES
serial*
            up 3-00:00:00
                                               NO
                                                         all
                                  1 no
1 no
serial*
                                                         all
            up 3-00:00:00
                                               NO
            up 3-00:00:00
serial*
                                                                593
serial*
            up 3-00:00:00
                                  1
                                               NO
                                                         all
                                                                226
parallel
            up 3-00:00:00
                               1-28
                                               NO
                                                         all
                                                                 1
parallel
            up 3-00:00:00
                               1-28
                                      no
                                               NO
                                                         all
                                                                101
            up 3-00:00:00
parallel
                               1-28
                                                         all
                                                                593 allocated c[3-24,26-75,...
                                      no
                                               NO
parallel
            up 3-00:00:00
                               1-28
                                                         all
                                                                226
                                               NO
                                      no
longrun
            up 14-00:00:0
                                                         all
                                      no
            up 14-00:00:0
                                  1
                                                         all
                                                                101
longrun
                                               NO
            up 14-00:00:0
                                 1
                                               NO
                                                         all
                                                                587 allocated c[3-24,26-75,...
longrun
            up 14-00:00:0
                                      no
                                               NO
                                                         all
                                                                226
                                                                          idle c[1-2,984-985]
                                1-2
test
            up
                    30:00
                                      no
                                               NO
                                                         all
            up 7-00:00:00
                                                         all
                                                                          mixed c[577-5781
hugemem
                                  1
                                      no
                                               NO
```

#### #SBATCH --constraint=snb

- The job is run only in Sandy Bridge (snb) nodes
- The other option is Haswell node (hsw) or
  - #SBATCH --constraint=hsw
- Either that is free "snb|hsw"
  - #SBATCH --constraint="snb|hsw"
- Currently the default is to use either architecture in serial and longrun partitions
- Sandy Bridge in test and parallel
- A single job cannot use CPUs from both architectures, but SLURM will take care of this

```
#!/bin/bash -1
#SBATCH -J myjob
#SBATCH -e myjob_err_%j
#SBATCH -o myjob_output_%j
#SBATCH --mail-type=END
#SBATCH --mail-
    user=a.user@foo.net
#SBATCH --mem-per-cpu=4000
#SBATCH -t 02:00:00
#SBATCH -n 1
#SBATCH -p serial
```

module load myprog
srun myprog -option1 -option2

#SBATCH --constraint=snb

# module load myprog srun myprog -option1 -option2

- Your commands
  - These define the actual job to performed: these commands are run on the compute node.
  - See application documentation for correct syntax
  - Some examples also from batch script wizard in SUI
- Remember to load modules if necessary
- By default the working directory is the directory where you submitted the job
  - If you include a cd command, make sure it points to correct directory
- Remember that input and output files should be in \$WRKDIR (or in some case \$TMPDIR)
- > \$TMPDIR contents are deleted after the job
- > srun tells your program which cores to use. There are also exceptions...



#!/bin/bash -1

#SBATCH -J myjob

#SBATCH -e myjob\_err\_%j

#SBATCH -o myjob\_output\_%j

#SBATCH --mail-type=END

#SBATCH --mailuser=a.user@foo.net

#SBATCH --mem-per-cpu=4000

#SBATCH -t 02:00:00

#SBATCH -n 1

#SBATCH -p serial

module load myprog
srun myprog -option1 -option2

#### Most commonly used sbatch options



#### Slurm option

--begin=time

-c, --cpus-per-task=ncpus

-d, --dependency=type:jobid

-e, --error=err

--ntasks-per-node=n

-J, --job-name=jobname

--mail-type=*type* 

--mail-user=user

-n, --ntasks=ntasks

-N, --nodes=N

-o, --output=*out* 

-t, --time=minutes

--mem-per-cpu=<number in MB>

--mem=<number in MB>

Description

defer job until HH:MM MM/DD/YY

number of cpus required per task

defer job until condition on jobid is satisfied

file for batch script's standard error

number of tasks per node

name of job

notify on state change: BEGIN, END, FAIL or ALL

who to send email notification for job state changes

number of tasks to run

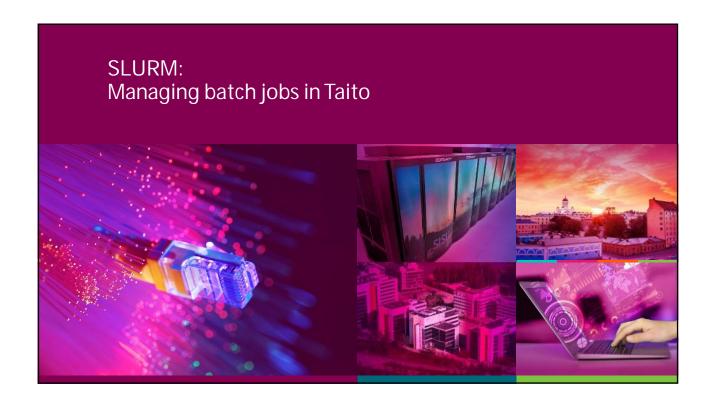
number of nodes on which to run

file for batch script's standard output

time limit in format hh:mm:ss

maximum amount of real memory per allocated cpu (core)

required by the job in megabytes maximum memory per node

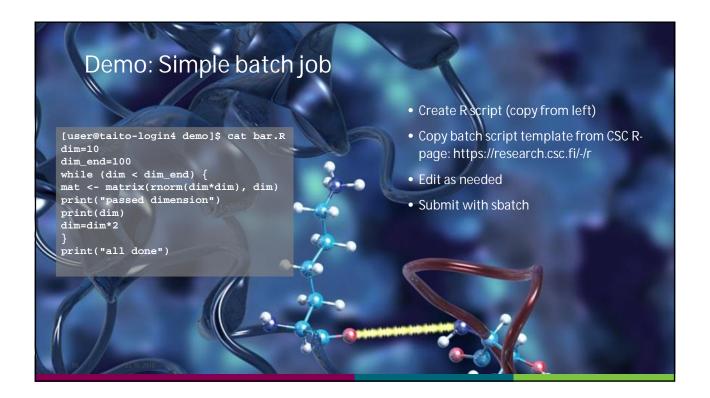


## Submitting, cancelling and stats of batch jobs



- The script file is submitted with command
  - \$ sbatch batch\_job.file
- Job can be deleted with command
  - \$ scancel <jobid>
- Display the used resources of a completed job
  - \$ seff <jobid>

10/23/2018



#### Queues



• The job can be followed with command squeue:

To estimate the start time of a job in queue

```
$ scontrol show job <jobid>
row "StartTime=..." gives an estimate on the job start-up time, e.g.
StartTime=2014-02-11T19:46:44 EndTime=Unknown
```

- scontrol will also show where your job is running
- If you add this to the end of your batch script, you'll get additional info to stdout about resource usage

seff \$SLURM\_JOBID

### Examples of seff outputs



[erkki@taito]\$ seff 52000797
Job ID: 52000797
Cluster: csc
User/Group: erkki/csc
State: COMPLETED (exit code 0)
Nodes: 4
Cores per node: 16
CPU Utilized: 00:37:22
CPU Efficiency: 87.58% of 00:42:40 corewalltime
Memory Utilized: 7.53 GB (estimated maximum)
Memory Efficiency: 3.21% of 234.38 GB
(58.59 GB/node)

Comments: only small part of memory used, could request less (now used the default 0.5GB/core), but for a parallel job like this, it's better to request full nodes anyway.

[erkki@taito]\$ seff 52000798\_6
Job ID: 52000798
Array Job ID: 52000798\_6
Cluster: csc
User/Group: erkki/csc
State: COMPLETED (exit code 0)
Nodes: 1
Cores per node: 4
CPU Utilized: 00:24:09
CPU Efficiency: 98.17% of 00:24:36
core-walltime
Memory Utilized: 23.50 MB
Memory Efficiency: 1.15% of 2.00 GB

Comments: only small part of memory used, could request less (now used the default 0.5GB/core). Theoretically the job used only 23.5/4=6MB/core, but asking for e.g. 100MB/core (for safety) would likely make the job queue less.

### Job logs



Command sacct can be used to study past jobs
 Useful when deciding proper resource requests

TIP: Check MaxRSS to see how much memory you need and avoid overbooking. See also command seff JOBID

\$ sacct Short format listing of jobs starting from

midnight today

\$ sacct -u <username> list only jobs submitted by username \$ sacct -o list only named data fields, e.g.

\$ sacct -o jobid,jobname,maxrss,reqmem,elapsed -j <jobid>

### Available Taito queues and limits



Queue	Maximum number of cores	Maximum run time	Maximum total memory
serial (default)	16 / 24 (one node*)	3 days	256 GB
parallel	448 / 672 (28 nodes*)	3 days	256 GB
longrun	16 / 24 (one node*)	14 days	256 GB
hugemem	40 (one node)	7 days	1.5 TB
test	32 / 48 (two nodes*)	30 min	64 GB

<sup>\*</sup> Sandy Bridge / Haswell (one Sandy Bridge node consists of 16 and Haswell one of 24 cores)

# Most frequently used SLURM commands

CSC

Command Description srun Run a parallel job.

sallocAllocate resources for interactive use.sbatchSubmit a job script to a queue.

**scancel** Cancel jobs or job steps.

**sinfo** View information about SLURM nodes and partitions.

squeue View information about jobs located in the SLURM scheduling queue smap Graphically view information about SLURM jobs, partitions, and set

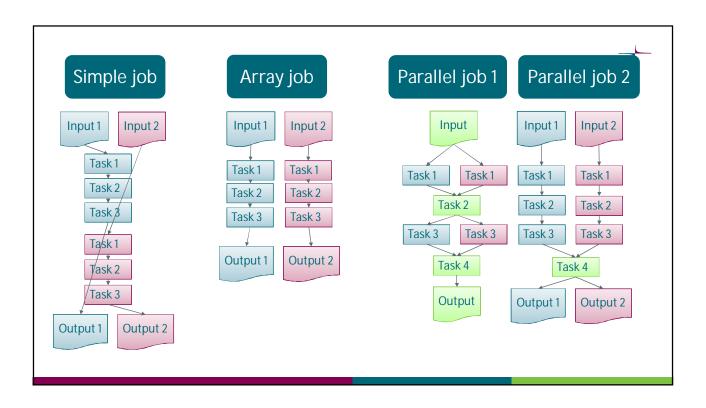
configurations parameters

**sjstat** Display statistics of jobs under control of SLURM (combines data from sinfo,

squeue and scontrol)

**scontrol** View SLURM configuration and state. **sacct** Displays accounting data for batch jobs.





# Simple job

- Pros: Easy, just check paths and packages.
- Cons: Slowest.

# Array job

- Pros: Relatively easy, just check paths and packages, and add using command line arguments. SLURM takes care of the job management
- Cons: Applicable only in certain cases

# Parallel job

- Pros: All kind of use cases can be done.
- Cons: Requires modifying code.
   More complicated code, because the job management is done in the code.



### Multi-core processing support in GIS-software



Most GIS software won't support parallel processing out of the box, but a few do:

- Taudem, SagaGIS partially
- In R and Python special packages for parallel processing
- (LasTools, ArcGIS Pro in Windows, but these can not be used in Taito.)
- Some others: https://research.csc.fi/geocomputing

# Array jobs for GIS



- Same analysis for:
  - different input files / map sheets
  - different scenarios
  - different variables
  - different time periods
- Suits well if the jobs are independent of each other, for example raster calculator.
- In many cases if using map sheets the borders need special care.

### Parallel jobs for GIS



- You have to divide the job to tasks:
  - o For vector data it might mean handling the data in chunks.
    - o For example, if calculating zonal statistics for 10 000 polygons, dividing them 1000 polygon chunks might work well.
  - o For raster data some kind of areal division is often most logical.
  - o For some repeated analysis it might be easy just to divide the queries to different tasks.
    - o For example, if routing or geocoding 1 000 000 addresses, dividing them 10 000 queries per task should work well.
- You normally have to test, what is a good size for data for one task. One task should take at least ca 10 min.

# Be careful with simultaneous opening of files



- If you are using the same input or output files in array or parallel jobs, depending on software it might cause trouble.
- Solutions:
  - o Make copies of input files for each processes you use.
  - o Write output to smaller files first, join them later

#### Array jobs



- #SBATCH --array=1-100
- Defines to run 100 jobs, where a variable
   \$SLURM\_ARRAY\_TASK\_ID gets each number (1,2,...100) in turn as its value. This is then used to launch the actual job (e.g.
- \$ srun myprog input\_\$SLURM\_ARRAY\_TASK\_ID > output\_ \$SLURM\_ARRAY\_TASK\_ID)
- Thus this would run 100 jobs:

```
srun myprog input_1 > output_1
srun myprog input_2 > output_2
...
srun myprog input_100 > output_100
```

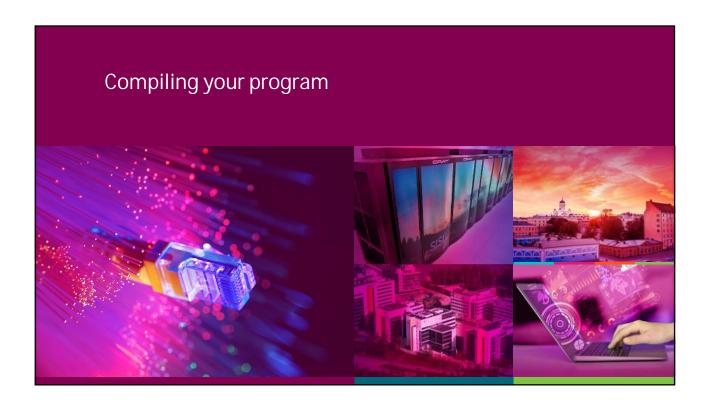
• For more information: research.csc.fi/taito-array-jobs

# Array jobs, output files



```
%j = job number#SBATCH -o array_job_out_%j.txt#SBATCH -e array_job_err_%j.txt
```

%A = ID of the array job and %a = \$SLURM\_ARRAY\_TASK\_ID
 #SBATCH -o array\_job\_out\_%A\_%a.txt
 #SBATCH -e array\_job\_err\_%A\_%a.txt



# What if the program you want to use is not installed?



- You can install it yourself
   Under your \$HOME, not in the "default" location/usr/bin)
- You can ask help for installation from <a href="mailto:servicedesk@csc.fi">servicedesk@csc.fi</a>
- A short overview:
  - $\circ \underline{\text{https://research.csc.fi/taito-installing-third-party-applications}}$

102

# How to compile a program?



- Typically the source code comes with instructions, read them.
- The CSC environments are explained at:

   https://research.csc.fi/taito-compiling-environment
   https://research.csc.fi/sisu-compiling-environment
- Always check that the code gives correct results

103

23.10.2018



#### R spatial

CSC

- R strong sides
  - Statistical analysis
  - Raster analysis
  - Time-series analysis
  - Point clouds
  - Faceted maps
  - Repeating workflows
  - Usage of external functions: SagaGIS, GRASS etc
  - Rstudio
  - Several packages for parallel computation

- R weak sides
  - Routing
  - 3D vector models
  - Interactiv map usage
  - Data editing
  - Different than GUI GIS software

### R in Taito, rspatial-env 1



This is an convinience module for loading easily spatial analysis related R tools. It loads the following software:

- module load rspatial-env
- <u>R</u> (3.5.0, 3.4.3, 3.4.1) with spatial packages:
  - ogeoR, geoRgIm, geosphere, ggmap, grid, gstat, GWmodel, lidR, mapproj, maptools, ncdf4, RandomFields, raster, rgdal, rgeos, rgrass7, rlas RSAGA, sf, sp, spacetime, spatial, spatial.tools, spatstat, spdep, strucchange.

### R in Taito, rspatial-env 2



- •rgdal GDAL/OGR (2.2.1), Proj4 (4.9.3)
- •rgeos GEOS (3.6.1)
- RSAGA\* Saga GIS (5.0.0)
- rgrass7 GRASS GIS (7.2.0, without GUI)
- \* RSAGA gives error messages: "Error: select a tool", but it seems to be <u>a bug of SAGA</u>. According to our tests it seems that RSAGA commands are working despite the message.
- $\hbox{* RSAGA io\_gdal library does not work, use gdal\_translate from gdalUtils.}$

# rspatial-env and Rstudio



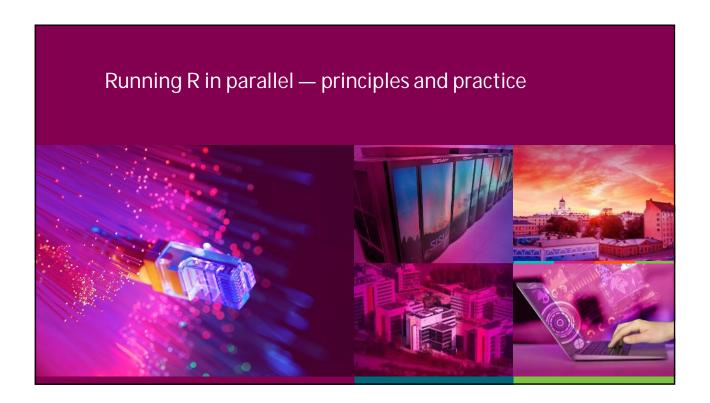
 In NoMachine use Rstudtio with GIS packages, use Applications -> Taitoshell -> GIS -> RStudio

### Installing R libraries



- Everybody can add R libraries for personal use
   These override the system ones
- Normally similar to local installations:

   install.packages("pkgname")
   or using Rstudio menus
- If the package needs some external library, the user installation is likely tricky.
- If you think that the package should be useful also for others or requires some external software to be installed, ask servicedesk@csc.fi



#### "My R code is slow... what can be done?"



- You should seek to have an understanding which part of your code is the time consuming one, and why, and how othe function system. time() can help

  - o understanding the details of the method you are using helps
  - o knowing how to run a smaller version of the problem helps a lot ounderstanding even the basics of time complexity helps
- Always be suspicious of for-loops!
- Having more memory *almost never* helps
- Going parallel may help

# Mere presence of more cores does nothing

- With respect to workload cores are not like locomotives but more like trucks: several cores can not work on the exact same task
- Simply taking your ordinary R code to a laptop with 8 instead of 1 otherwise similar cores gives *no speed* boost at all.
- Somebody (possibly you) has to do something to divide the workload



By Douglas W. Jones - Own work, Public Domain, https://commons.wikimedia.org/w/index.php?curid=7756596



https://commons.wikimedia.org/w/index.php?curid=228396

#### Parallel scaling



- Unfortunately, increasing the number of cores will *never* decrease the time needed in the same proportion
- Not all parts of the program will benefit from parallel computing (Amdahl's law)
- Parallel computing introduces extra work that wouldn't need to be done when computing serially (overhead)
- Due to these facts increasing the number of cores may help only up to a point but no further, or even not at all, and it may depend on many things such as data size

# Level zero: not even parallel, just simultaneous



- Can you imagine going around in a computer classroom, firing up R on every computer and letting each one work on part of your problem?
  - o such as each fitting the same model to 20 different data sets or with 20 different combinations of parameters
- Do you also have access to a cluster with R installed on it?
   Good! Then you can do the same without even getting up from your chair.
- Upside: really easy. Downside: limited usability

Array jobs in Taito

#### Array job with R, giving the mapsheet path as argument



```
#!/bin/bash
                                              module load rspatial-env
#SBATCH - J array_job
                                              # move to the directory where the data files locate
#SBATCH-o array_job_out_%j.txt
                                              cd ~/git/geocomputing/R/contours/array
#SBATCH-e array_job_err_%j.txt
                                              # set input file to be processed
#SBATCH-t 00:02:00
                                              name=$(sed -n "$SLURM_ARRAY_TASK_ID"p../mapsheets.txt)
#SBATCH -- mem-per-cpu=4000
                                              # run the analysis command
#SBATCH -- array=1-3
                                              srun Rscript Calc_contours.R$name
#SBATCH-n1
#SBATCH-p serial
```

# Array jobs with R, reading the argument in R script



```
args = commandArgs(trailingOnly=TRUE)

if (length(args)==0) {
  stop("Please give the map sheet number", call.=FALSE)
} else if (length(args)==1) {
  mapsheet <- args[1]
}</pre>
```

# Running R code in parallel

- -

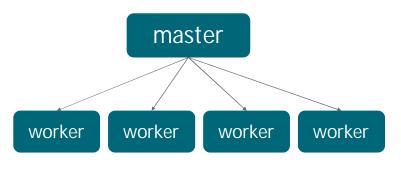
- There are several R packages for parallel computing:

   parallel (snow)
   oforeach and doMPI
  - o Rmpi

# snow (parallel)



• This R package offers support for simple parallel computing in R, following the master - workers paradigm



### parallel (snow), batch job file



The execution command that needs to be added to the end of the batch job file given above is like this:

 $srun\,RMPISNOW\,\hbox{--no-save-f myrscript}. R$ 

# parallel (snow), R script



The script should contain a getMPlcluster call which will produce the reference to the cluster that can be given to various other functions, like in this example:

```
cl<-getMPIcluster()
funtorun<-function(k) {
   system.time(sort(runif(1e7)))
}
system.time(a<-clusterApply(cl,1:7,funtorun))
a
stopCluster(cl)</pre>
```

Only the master process continues to run the given script.

#### foreach and doParallel



- The foreach package implements a for-loop that uses iterators, and also allows for parallel execution using its %dopar% operator.
  - o now it is definitely you who needs to decide how the work gets divided, or at least one aspect of that
- It comes with several "parallel backends", of which the doMPI package should be used on Taito.
- From the foreach vignette: "Running many tiny tasks in parallel will usually take more time to execute than running them sequentially"
  - o so now you need to start paying attention to details

# foreach and doMPI, batch job file



The execution command that needs to be added to the end of the batch job file given above is like this:

srun Rscript --no-save --slave myrscript.R

Note that this starts a number of R sessions equal to the number of reserved cores that all start to execute the given script. The --slave option prevents all of them printing the R welcome message etc. to the common output file.

# foreach and doMPI, R script



The script should include a call to startMPIcluster very close to the beginning, as all the processes will execute everything before that call, while only the master continues after that call.

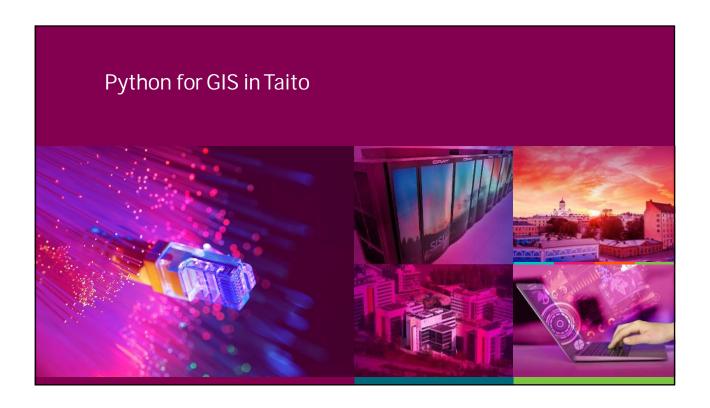
```
library(doMPI,quietly = TRUE)
cl<-startMPIcluster()
registerDoMPI(cl)
system.time(a<-foreach(i=1:7) %dopar% system.time(sort(runif(1e7))))
a
closeCluster(cl)</pre>
```

#### Further information



- CSC R documentation: https://research.csc.fi/-/r
- rspatial-env module: https://research.csc.fi/-/rspatial-env

Support: servicedesk@csc.fi



### Geoconda

- NumPy, Scipy, Pandas etc.
- GIS Specific packages
  - Geopandas
  - Fiona
  - Shapely
  - Rasterio
  - Rasterstats
  - cartopy
  - GDAL/OGR
  - Networkx
  - Skimage
  - Pyproj
  - Pysal
  - Rtree
  - Descartes

module load geoconda

126



#### Installing own Python packages



- Python packages not available by CSC can be installed by user
- Easy installation from Python Packaging Index
  - opip install -user package\_name
  - o Package is installed under \$HOME/.local and is available without further actions
  - o Pip tries to take care also of dependencies
- Installing from source with distutils opython setup.py install - -user
- Alternate installation directories
  - o Both pip and distutils have -prefix option for specifying installation directory
  - opip install --prefix=\$HOME/my\_modules package\_name
  - o Directory has to be added to PYTHONPATH:
  - o export PYTHONPATH=\$PYTHONPATH:\$HOME/my\_modules/lib/python2.7/site-packages

127



### Creating your own conda environment

- Create new environment
  - conda create -n my\_tools
- Activate/deactivate your new environment
  - source activate my\_tools
  - source activate my\_tools
- Install new packages
  - conda install -n my\_tools geopandas
- https://conda.io/docs/user-guide/tasks/manage-environments.html#

128

#### Running Python code in Taito and Taito Shell



- Same as most other software
- In Taito Shell: python your\_script.py args
- To submit batch jobs: sbatch batch\_job.sh

```
#!/bin/bash -1
#SBATCH -J python_focal
#SBATCH -o out.txt
#SBATCH -e err.txt
#SBATCH -t 00:00:20
#SBATCH --cpus-per-task=1
#SBATCH --mem-per-cpu=2
#SBATCH -p serial

# load needed modules
module load geoconda
srun python your_script.py args
```

# Array jobs and python



· Read arguments from file row by row and pass to python script

```
#!/bin/bash -1 Batch job file

#SBATCH -J python_focal

#SBATCH -o array_job_out_%A_%a.txt

#SBATCH -e array_job_err_%A_%a.txt

#SBATCH -t 00:00:20

#SBATCH --cpus-per-task=1

#SBATCH --mem-per-cpu=2

#SBATCH -p serial

#SBATCH --array=1-3

# load needed modules

module load geoconda

name=$(sed -n "$SLURM_ARRAY_TASK_ID"p file_names.txt)

srun python your_script.py $name
```

```
your_script.py
import sys
filename = sys.argv[1]
...do something with the file..
```

### Parallel jobs with Multiprocessing Python module



import multiprocessing as mp from time import sleep

def worker(x):
 sleep(0.5)
 return x

if \_\_name\_\_ =='\_\_main\_\_':
 pool = mp.Pool(processes=4)
 print pool.map(worker, range(4))

```
from time import sleep

def worker(x):
    sleep(0.5)
    return x

if __name__ == '__main__':
    print [worker(i) for i in range(4)]
```

#### cProfile



- Tool to help you find out which part of your porgram takes most execution time
- Let's say function A takes 99% execution time
  - There's little point trying to optimize performance of anything else even if it is very inefficient.
- Usage:
  - python -m cProfile my\_python\_script.py

#### Further information



- CSC Python documentation: research.csc.fi/-/python
- Geoconda module: https://research.csc.fi/-/geoconda
- Geo-env module: https://research.csc.fi/-/geo-env
- Multiprocessing: https://docs.python.org/3/library/multiprocessing.html
- cProfile https://docs.python.org/2/library/profile.html

Support: servicedesk@csc.fi

133



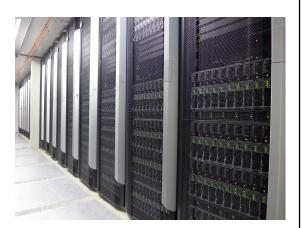
### Taito / Taito-shell pre-installed software for GIS



- $\circ R$
- o Python
- oMatLab/Octave
- oGDAL/OGR
- o GRASS GIS
- o LasTools (some)
- oPDAL
- o Proj4
- o QGIS
- o SagaGIS
- $\circ\, Taudem$
- o Zonation

 $\underline{https://research.csc.fi/software} {\:\raisebox{3pt}{$->$}} \ {\sf Geosciences}$ 

8.0.2



# GIS Software not available in Taito



#### Windows software:

- ArcGIS
- MapInfo
- LasTools Windows tools

#### Server software

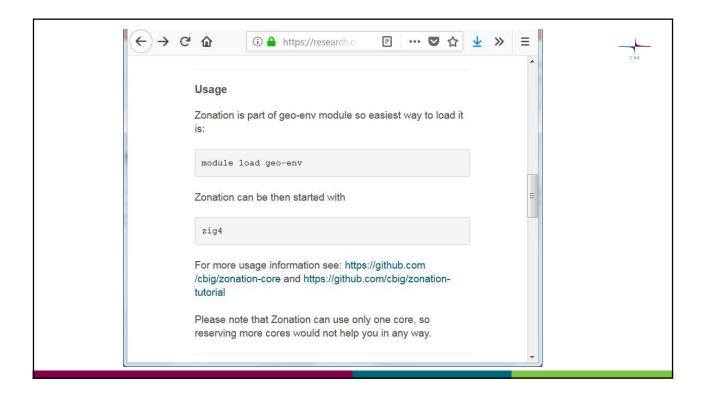
- GeoServer, MapServer
- PostGIS

### Web map libraries

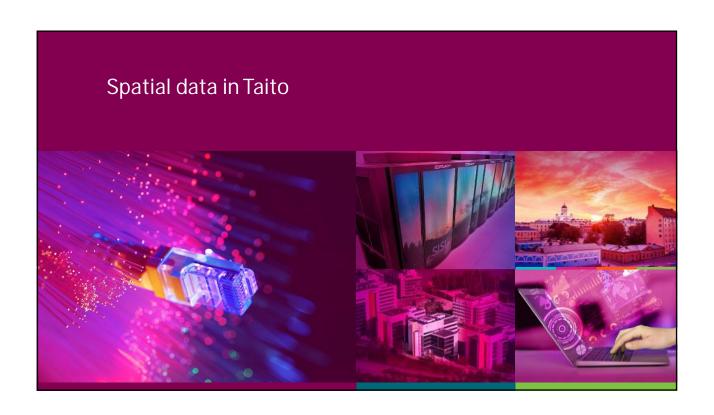
· OpenLayers, Leaflet

136





#### Using different GIS-software in Taito Python Bash QGIS GDAL Χ Х Χ Χ GRASS Χ (x) Х Χ (x) LasTools (x) Х Χ SagaGIS Χ (x) Χ Χ Taudem Х (x) (x) R spatial packages Χ Python geo packages



#### Shared data area in Taito

- All Paituli open data
- LUKE

Multi-source national forest inventory

NLS

Virtual rasters for DEMs

- Hosts large commonly used datasets
- Reduces the need to transfer data to Taito
- Located at /proj/ogiir-csc/
- All Taito users have read access.
- Only CSC personnel have write access.
- For data with open license
- If you think some other dataset should be included here, ask from servicedesk@csc.fi

https://research.csc.fi/gis\_data\_in\_taito

141

#### Virtual rasters

- csc
- Allows working with dataset of multiple files as if they were a single file.
- XML pointing to actual raster files
  - The virtual file doesn't need to be rectangular, it can have holes and the source files can even have different resolutions
- Taito has ready made virtual rasters for elevation models and a python tool to create your own for a specific area.

142

#### Virtual Rasters in Taito



- Ready made virtual rasters for 2m and 10m dems
- Corresponding to folder structure
- External overviews and xml headers

•

- A python script to create your own for a specific area
- Syntax:
- python/proj/ogiir-csc/mml/karttalehtijako/vrt\_creator.py [-h] [-i] [-o] [-p prefix] dataset < dem2m/dem10m > polygon output\_dir
  - -h print help
  - -i create individual vrt file for each polygon in shp
  - -o create overviews
  - -p refix> adds a prefix to output files

•

#### Virtual Rasters with different software



- Can be read by anything that uses GDAL drivers:
- QGIS, GRASS, SAGA, TAUDEM, Python and R packages, GDAL itself
- Virtual rasters can be of different size, working with big virtual rasters does not suit all.
  - QGIS is the best in vizualizing virtual rasters (if overviews are available).
  - Reading a smaller area of a big virtual raster for a certain analysis works well in Python rasterio, R raster packages and GRASS.
- But there are limitations...
  - GRASS requires linking with r.external, but still there is no need to create copies of actual data
  - SAGA uses gdal only for importing data so you can create a .sgrd copy of data, not use vrt directly.
- A lot of tools output raster of similar dimensions as input and currently nothing will create a tiled virtual raster as output.

#### Further information



- Geocomputing general: https://research.csc.fi/geocomputing
- GIS data in Taito: https://research.csc.fi/gis\_data\_in\_taito
- Virtual rasters: https://research.csc.fi/virtual\_rasters

Support: servicedesk@csc.fi

145



## CSC's cloud services and cPouta, setting up a virtual machine

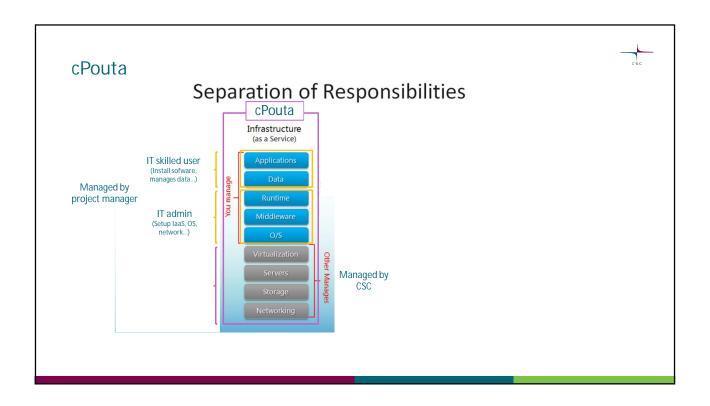


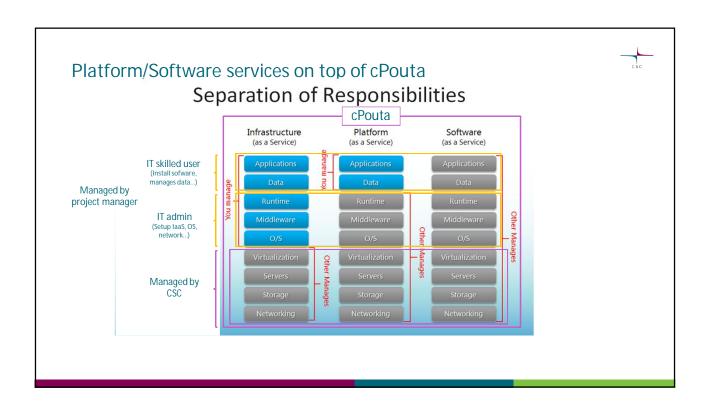
- Part 1: Cloud brainwashing
  - o Cloud concepts
  - o cPouta web interface
- Part 2: setting up a virtual machine
  - o Pouta concepts
  - o Virtual machine's settings

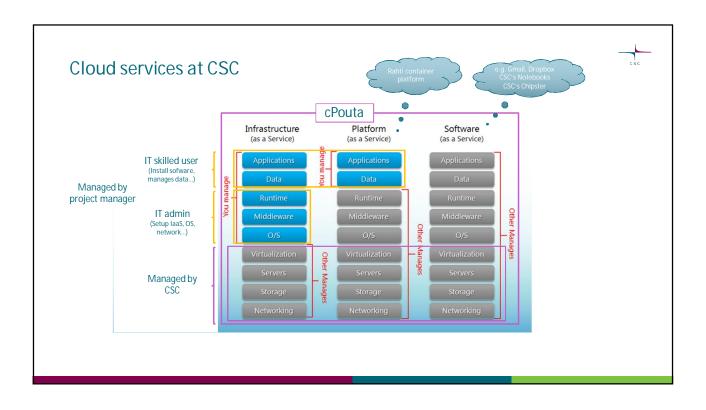
#### What cloud?

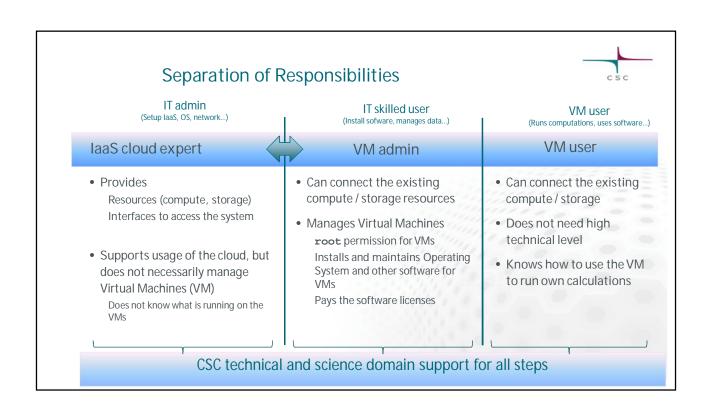


- Terminology overload, cloud used to mean e.g.:
  - Storage services (Dropbox)
  - o Virtual server hosting (Amazon Web Services)
  - o Software platforms (Google App Engine)
  - o Pretty much any web service (gmail, MS Office 365, ArcGIS Online)
  - o The Internet as a whole
- Self-service and automation are the common features









# Traditional HPC (Taito) vs. IaaS (cPouta)

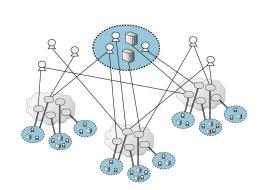


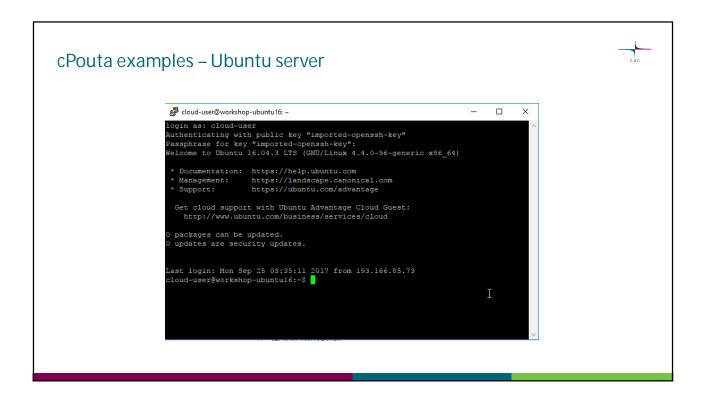
	Traditional HPC environment	Cloud environment Virtual Machine	
Operating system	Same for all: CSC's cluster OS	Chosen by the user	
Software installation	Done by cluster administrators Customers can only install software to their own directories, no administrative rights	Installed by the user The user has admin rights	
User accounts	Managed by CSC's user administrator	Managed by the user	
Security e.g. software patches	CSC administrators manage the common software and the OS	User has more responsibility: e.g. patching of running machines	
Running jobs	Jobs need to be sent via the cluster's Batch Scheduling System (BSS = SLURM in Taito)	The user is free to use or not use a BSS	
Environment changes	Changes to SW (libraries, compilers) happen.	The user can decide on versions.	
Snapshot of the environment	Not possible	Can save as a Virtual Machine image	
Performance	Performs well for a variety of tasks	Very small virtualization overhead for most tasks, heavily I/O bound and MPI tasks affected more	

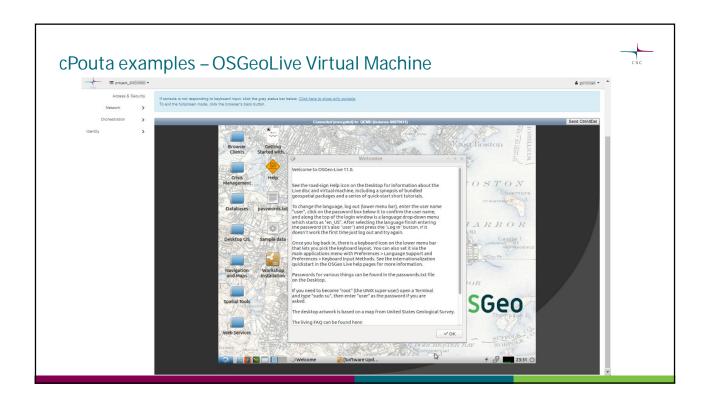
#### cPouta use cases

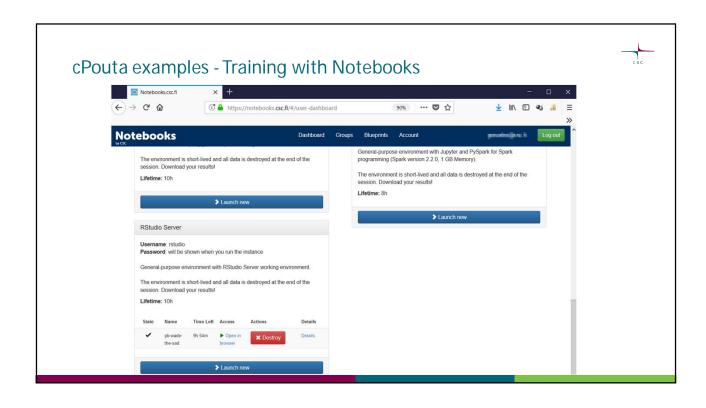


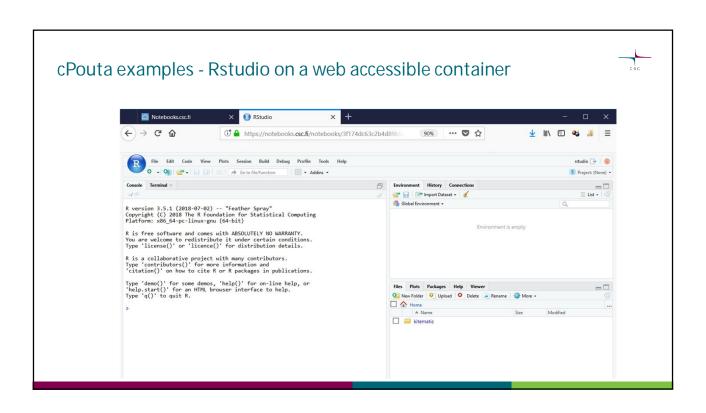
- Running scientific applications
  - o Computational clusters
    o Pre/post process Sisu/Taito bound data
- Running other types of software stacks
   Containerized or non-containerized
   Web/file servers, load balancers, databases etc
- Virtual computer class
- Ad hoc research data/information sharing

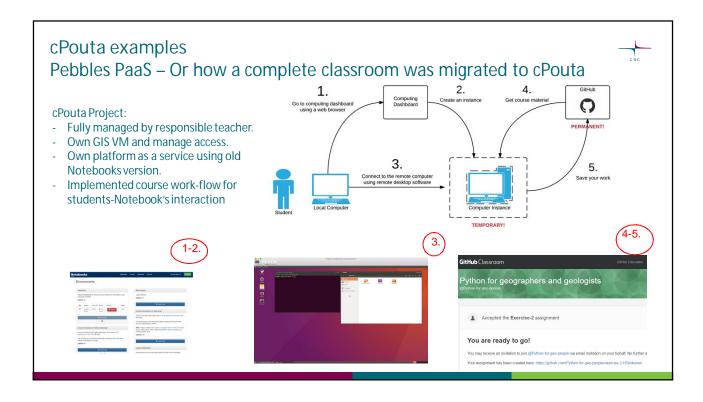


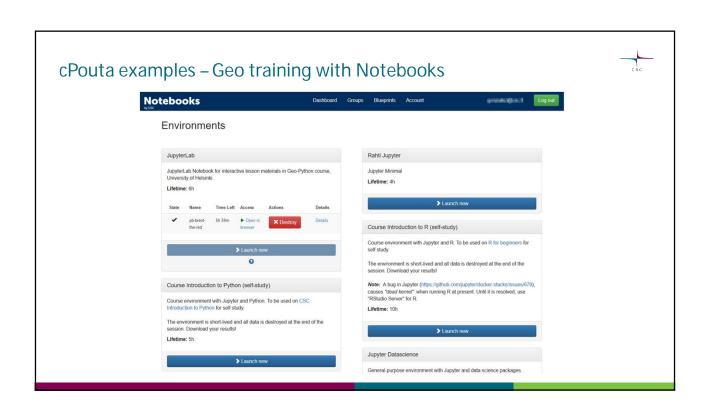


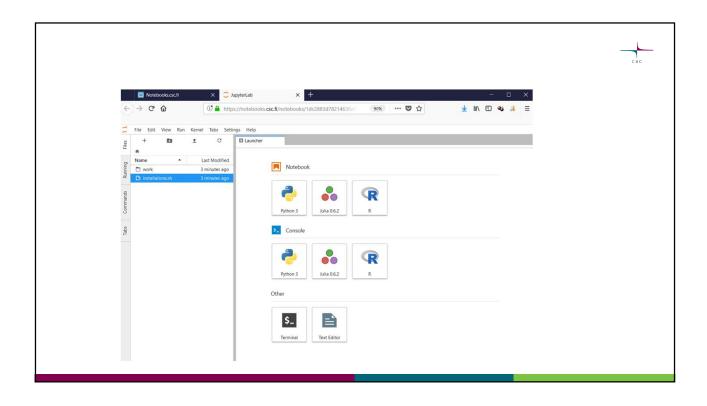


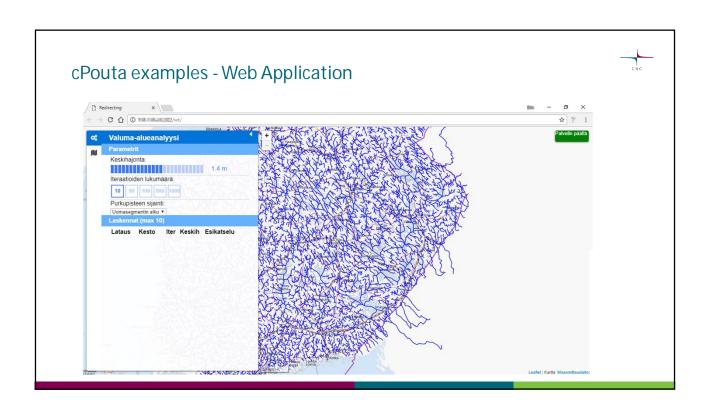












#### GIS related services



- Web map services
  - o GeoServer
  - o MapServer
- Databases
  - o PostgreSQL/PostGIS
- ArcPy
- Windows\* software

## Object storage



- Used for storing and sharing data / files.
- Included in Pouta projects
- Ready to use (no need to set up a virtual machine)
- Manage via the Pouta Web interface or via API (s3, swift, python...)
- Data can be accessed from anywhere using URL or via API
- Data can be private, public and temporarily shared
- Limitations:
  - o Object storage file can not be edited (you can delete and make a new copy)
  - o Not suitable for databases
  - o Can not (efficiently) be mounted as file system

164

8.6.2017

#### Rahti



- Platform-as-a-service (based on OpenShift, Red Hat's distribution of Kubernetes)
- Used for running and orchestrating containers that run applications
- Still you need to install your software and pack it as containers
- Same end goal as cPouta: enable end users to run their own software in the cloud oweb applications
  - o APIs/microservices for science
  - o Apache Spark
  - o Jupyter notebooks
- Compared to Pouta, you don't need to manage virtual machines but you need to manage containers

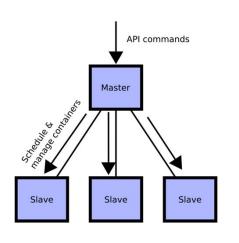
165

8.6.2017

## Advanced cloud applications - Container orchestration



- Running docker containers with Kubernetes
- o Using Magnum for creating resources
  - o Used to dividing work between different cloud machines
  - ohttps://research.csc.fi/pouta-container-orchestration
- Use cases of cloud geocomputing with workload management framework:
  - o JRC Earth Observation Data and Processing Platform
  - o Über
  - o EOX-Sentinel2 mosaic
  - o Mapillary



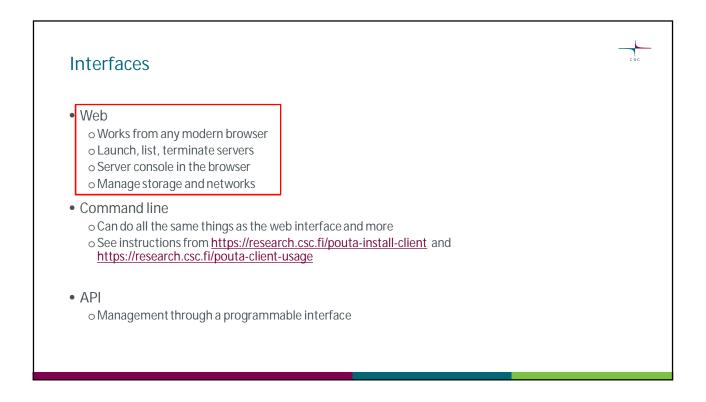
#### Pouta Clouds in general

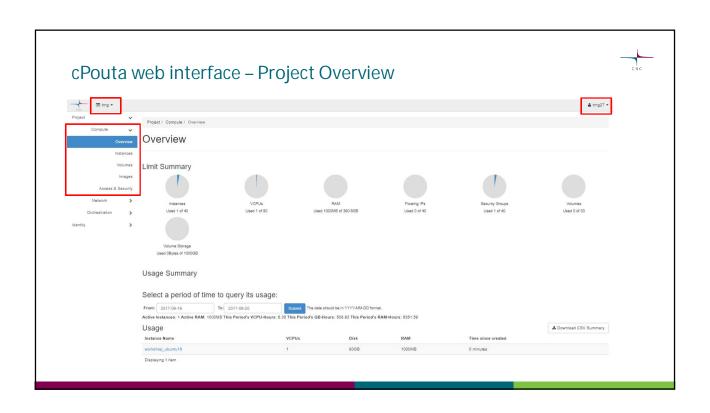


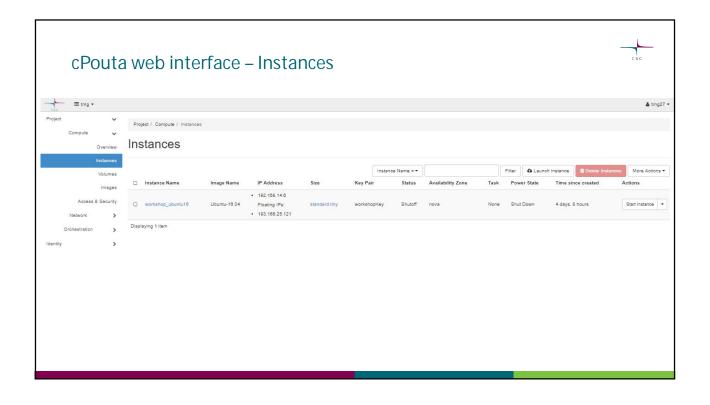
- Serviced offered by CSC (hardware in Finland)
- True self-service cloud I aaS powered by OpenStack
  - Deploy your own virtual machines, storage and networks as your requirements evolve
  - o No proprietary software to limit scalability
- Simple to create and modify virtual resources
   Choose from Web UI, CLI or RESTful APIs
- Designed to serve scientific as well as other use cases
  - o General purpose
  - o High Performance Computing
  - o Data Intensive Computing
  - o Sensitive data

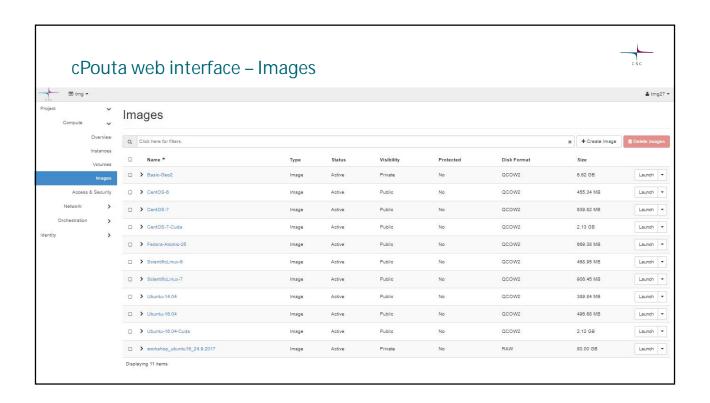


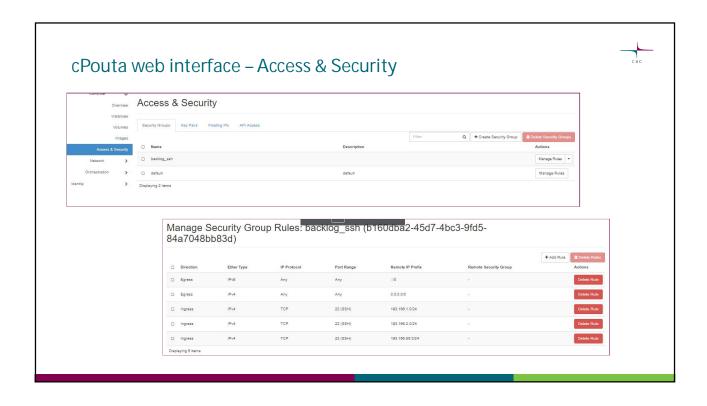
#### **Pouta Clouds** cPouta Cloud ePouta Cloud Public type service "Virtual Private Cloud" General description In production since 2013 Direct connections to customer networks From general pupose use to specialized Designed specifically for Use cases scientific computing handling sensitive data ISO/IEC27001 compliant Yes Yes VAHTI 2/2010 raised information security Yes Yes requirements Min. Edu. funded Yes Yes for open Finnish research and education **OPN/MPLS** VM access Internet VM installation, Firewall configuration, Load Self-service Self-service Balancing, VM auto-recovery, Backups No limits No limits **Supported Operating Systems** For more information: https://research.csc.fi/cloud-computing

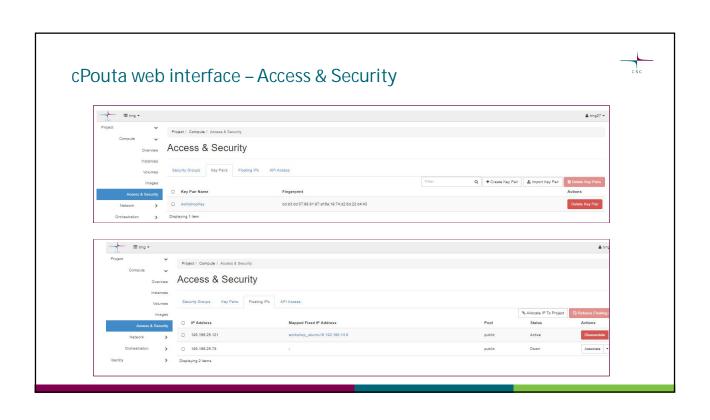


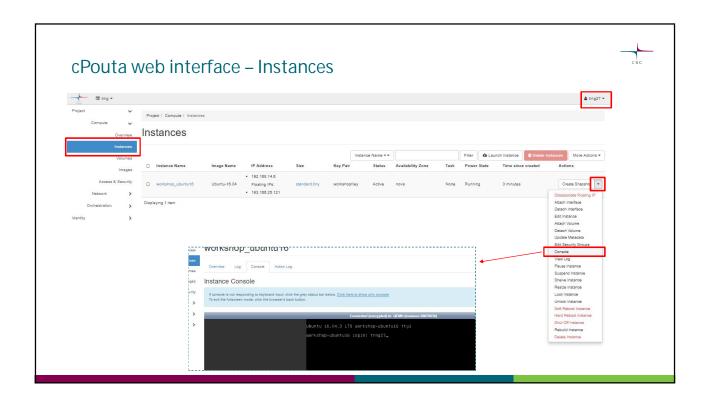


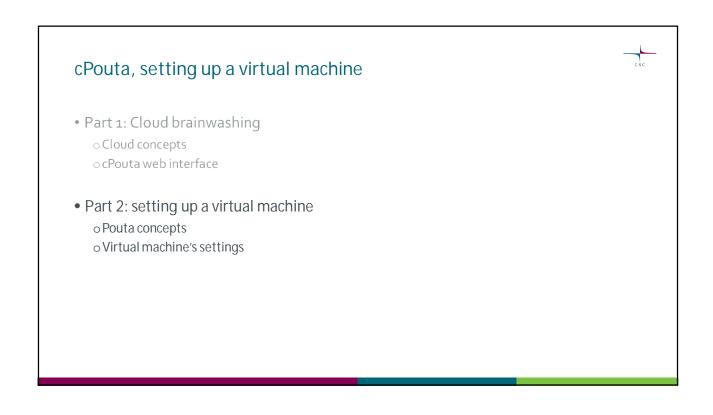












## Options for installing software



- From an image or Docker container with ready installed software packages
  - ofor ex. OSGeoLive
- Installing sofware manually
   ofor ex. using apt-get command line
- Scripting tools ofor ex. Ansible

## Virtual Machine Images



- Images are templates for launching (creating) VMs
  - o You can also launch VMs from volume snapshots (an image created from a VM or a bootable volume)
- CSC provides ready made images
  - o Based on different OSs
  - o Updated regularly
- Own images could be uploaded for own project
  - o For ex. created with VirtualBox or Vmware
  - o You can also use installation media to create a VM from scratch and then create an image out of it

#### Virtual Machines



- Virtual Machine = VM = Instance
  - o A virtual server managed by a OpenStack user
  - o User normally logins with ssh key pair
  - o Console access via OpenStack Web UI or ssh connection
  - o User gets root permissions and thus can create user accounts independent of the cloud middleware user accounts.
  - $\circ\,\text{VMs}$  are launched using an image

#### Virtual Machines - Flavors in cPouta



Flavors specify the resources of a VM
 CPU cores, RAM, root disk, ephemeral disk

Flavor	Cores	Memory/core	Memory	Disk (root)	Disk (ephem.)
standard.small	2	1 000 MB	2 000 MB	80 GB	0 GB
hpc-gen1.4core	4	3 750 MB	15 000 MB	80 GB	0 GB
hpc-gen2.8core	8	5 000 MB	40 000 MB	80 GB	0 GB
Io.70GB	2	5 000 MB	10 000 MB	20 GB	70 GB
gpu.1.1gpu	1 GPU (14 cores)	8 928 MB	125 000 MB	80 GB	0 GB

For more information: https://research.csc.fi/pouta-flavours

#### Virtual Machines - Flavors in cPouta



VM type	Description	Use	
standard.*	Oversubscribed "traditional" cloud virtual machines.	All non-CPU, non-IO intensive workloads	
hpcgen1.*	Non-oversubscribed, non-HT Sandy Bridge nodes (Taito)	CPU intensive HPC/HTC workloads	
hpcgen2.*	Non-oversubscribed, HT Haswell large memory nodes	Memory and CPU intensive HPC/HTC workloads	
io.*	SSD-backed high IOPS nodes	IOPS intensive workloads	
gpu.*	SSD-backed NVIDIA Tesla P100 GPGPU	HPC applications leveraging GPUs Machine- and deep learning Rendering	

For more information:  $\underline{\text{https://research.csc.fi/pouta-flavours}}$ 

# Volumes (storage types)



- Root disk
- Ephemeral storage
- Persistent volumes
- Volume snapshots
   o (Volumes can also be stored as Images)

No back-up by CSC!

#### Access & Security - Virtual Machines and Internet



- New VMs are not connected to Internet by default
- You need to allocate an IP address to it
- Once connected to Internet anyone could access/use it
- You are responsible of your VMs security
  - o Operating system, applications, user accounts, firewalls...
  - o SSH keypairs
  - o Security groups

# Access & Security - Floating IPs (= Public IPs)



- Access to VMs from Internet is provided by floating IPs
- You need to assign IPs to specific VMs
- You can disassociate IP from a VM and it goes back to the pool
- You can allocate IPs for your project (pool of IPS)
  - o Allocated IPs are ready to be used by VMs
  - o These IPs stay the same until you release them

## Access & Security - SSH Key Pairs



- SSH key pairs are used to secure login to the VMs
  - o You only need to create it once for each user
  - o New VMs created from CSC provided images can only be accessed with a SSH key (you can change this later if necessary)

For more information: <a href="https://research.csc.fi/pouta-getting-started">https://research.csc.fi/pouta-getting-started</a>

# Access & Security - Security Groups



- Security Groups are sets of firewall rules which limit access to your VMs
  - o By default, all the ports are closed
  - o Avoid adding rules to the "default" rule set
  - o An instance can have several security groups
  - o You can edit/create security groups at any time
  - o You can add/remove security groups to a VM at any time
  - o Keep as closed as possible i.e. limit the IP range if possible

For more information: <a href="https://research.csc.fi/pouta-getting-started">https://research.csc.fi/pouta-getting-started</a>