#### 1. SLURM: introduction

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#### Limited resources

- Cluster has many users wanting to run jobs, which limits:
  - CPU
  - Working memory
  - Time
- How to assign which resources to which job?

# Job scheduling

#### Job (computing)

From Wikipedia, the free encyclopedia

In computing, a job is a unit of work or unit of execution (that performs said work).

#### Job scheduler

From Wikipedia, the free encyclopedia

A job scheduler is a computer application for controlling unattended background program execution of jobs.[1]

#### **SLURM**

- Simple Linux Utility for Resource
   Management
- Job scheduler on:
  - UBELIX
  - IBU cluster



# Resource allocation commands

- sbatch
- srun
- salloc

```
#!/usr/bin/env bash

my_program \
--cpu 32 \
--memory 128G
```

\$ sbatch --cpus-per-task=32 --mem-per-cpu=4G ./script.sh

# #!/usr/bin/env bash #SBATCH --cpus-per-task=32 #SBATCH --mem-per-cpu=4G my\_program \ --cpu 32 \ --memory 128G

\$ sbatch ./script.sh

# 2. Frequently used sbatch options

# 2.1 user specific

- Job name:
  - --job-name=my\_job\_name
- e-mail
  - --mail-user=user@students.unibe.ch
  - --mail-type=begin,end,fail

#### 2.2 output & error

- --output=existing/path/output\_%j.txt
- --error=existing/path/error\_%j.txt

Path should exist!
Job will fail otherwise
(without error message)

# 2.3 Required resources

- CPU
  - --cpus-per-task=2
- Working memory
  - --mem-per-cpu=4G
- Time (days-hours:minutes:seconds)
  - --time=1-05:00:00

Conservative values will cause your job to start earlier **But:** job will fail if resources are overrequested!

# 3. Interactive jobs

# Why submit interactive job?

- Interactive job: allocated resources that are approachable with shell
- Head (login) node is not for computation
- Debugging and testing can be much more convenient if interactive

#### srun

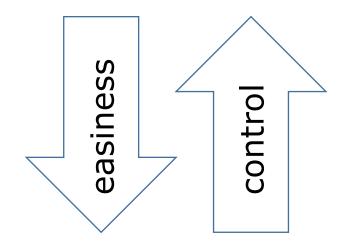
- Versatile command
- Used for job steps within sbatch (not treated in this course)
- Also for allocation of interactive job with pty (pseudo-terminal mode)

```
$ srun --cpus-per-task=1 --mem-per-cpu=4000 \
> --time=00:05:00 --pty bash
```

# 4. Modules

#### Software

- Install it yourself (at ~)
- Use a container
- Install with conda
- Use modules



#### Modules

- Check for available modules:
  - module avail
- Load a module in environment:
  - module load
- Unload a module:
  - module rm
- Available modules:
  - https://www.vital-it.ch/services

# 5. Job arrays

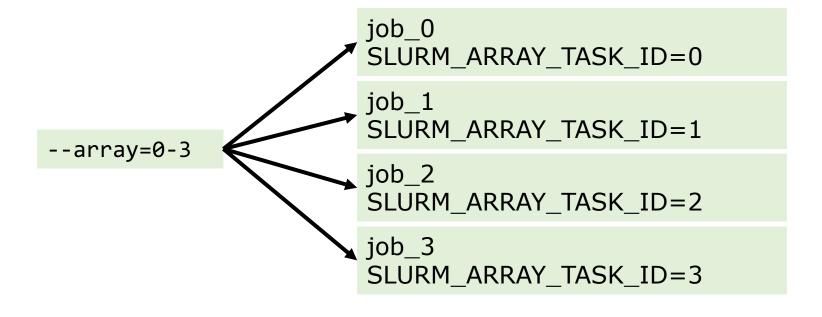
# 5.1 Jobs in parallel

- Run similar command with different parameters: parameter sweep
- E.g. alignment (e.g. with minimap2) on several files.

```
#!/usr/bin/env bash

#SBATCH --array=0-3
#SBATCH --cpus-per-task=32
#SBATCH --mem-per-cpu=4G

my_program \
$SLURM_ARRAY_TASK_ID
```



# 5.2 Using UNIX arrays

```
$ 1s
file1.txt file2.txt file3.txt
$ FILES=(./*)
$ echo ${FILES[0]}
                           UNIX uses zero-
file1.txt
                            based indexing
$ echo ${FILES[1]}
file2.txt
$ echo ${FILES[2]}
file3.txt
```

#### script.sh

```
#!/bin/bash

#SBATCH --cpus-per-task=32

#SBATCH --mem-per-cpu=4G

#SBATCH --array=0-7

FILES=(/path/to/input_data/*)

my_program ${FILES[$SLURM_ARRAY_TASK_ID]}
```