



Hands-on training for PIConGPU users

getting started with PIConGPU on LUMI

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Introduction

What will we do today?

- check your access to LUMI
- setup and run first PIConGPU simulation
- data analysis of simulation using openPMD and Jupyter
- dive deeper into how to setup a PIConGPU simulation
- **5** time for questions
- start another project on your own







Part I

Check LUMI access

Checking you are part of the workshop project

- go to: https://my.lumi-supercomputer.eu/
- login with your credentials
- go to projects and check whether you have access to ENCCS training (EHPC-DEV-2024D09-003)







Setting up an SSH key pair

create a (LUMI specific) key pair:

```
ssh-keygen -t ed25519 -f /home/username/.ssh/id_juwels_ed25519
```

- now you should have:
 - your private key: id_lumi_ed25519
 - and your public key: id_lumi_ed25519.pub
- go to: https://mms.myaccessid.org/profile/

MENU

* Settings

* My profile

* My linked accounts

* Settings







ssh into LUMI

execute:

ssh -i /.ssh/id_lumi_ed25519 your_username@lumi.csc.fi

you should see:







vour workshop project

check which projects you have access to:

lumi-allocations Data updated: 2024-10-21_17:03:48 CPU (used/allocated)| Project

113/512000

project_465001310 compare to:



24/18000

(0.0%) core/hours

(0.0%) core/hours

GPU (used/allocated)|

(0.0%) gpu/hours|

(0.1%) gpu/hours



Storage (used/allocated)

145/90000

(0.0%) TB/hours

(0.2%) TB/hours



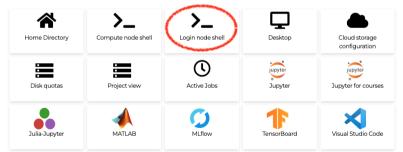
alternative access via browser

■ LUMI offers a web interface

www.lumi.csc.fi

- login with you credentials
- select compute node shell

Pinned Apps









creating your working directories

we are working with

project_465001310

project directory

/projappl/project_465001310

- please create a directory there
 cd /projappl/project_465001310
 mkdir \$(whoami)
- simulation directory

/scratch/project_465001310

please create a directory there
cd /scratch/project_465001310
mkdir \$(whoami)









Part II

First PIConGPU simulation

There is a manual!

https://picongpu.readthedocs.io

is your first source of help!

The steps we follow now are similar to the 'PIConGPU in 5 Minutes on Hemera' tutorial there.







Performing PIConGPU simulations

The general workflow

- Get PIConGPU
 \$ git clone https://github.com/ComputationalRadiationPhysics/picongpu.git
- **2** Prepare and load environment
 - \$ source picongpu.profile
- 3 Create simulation setup
 - \$ pic-create \$PIC_EXAMPLES/LaserWakefield myLWFA
- Adjust setup parameters
 - \$ vim simulation.param
- **5** Compile setup
 - \$ pic-build
- 6 Run setup (submit to batch system)
 - \$ tbg -s -t -c etc/picongpu/*.cfg mySimOutputDirectory
- Analyze results
- 12/35 · \$ jupyter-lab

(1) get the source code

download the PIConGPU source code:

```
in home directory
cd $HOME/
mkdir src
cd src
```

get source code

```
git clone https://github.com/ComputationalRadiationPhysics/picongpu.git
```





- (2) prepare and load environment
 - copy default setup file:

```
cp picongpu/etc/picongpu/lumi-eurohpc/lumi-G_hipcc_picongpu.profile.example
$HOME/lumi-G_hipcc_picongpu.profile
```

- either use your editor of choice in the terminal (vim, emacs, nano) or use the web interface
- adjust the following lines:
 - 8 export MY_MAILNOTIFY="ALL"
 - 9 export MY_MAIL="your.mail@server.gov"
 - 16 export PROJID=project_465001310
 - 17 export PROJECT_DIR=/projappl/\$PROJID/
 - 18 export SCRATCH=/scratch/\$PROJID/
 - 27 export EDITOR="emacs -nw"
 - 72 export PIC_LIBS=\$PROJECT_DIR/workshop_software/local
 - 81 export PICSRC=\$HOME/src/picongpu

source \$HOME/lumi-G_hipcc_picongpu.profile





(3) Create simulation setup

```
cd $HOME
mkdir -p picInputs
cd picInputs
pic-create LaserWakefield LWFA
cd LWFA
```

- ..
- peek into include/picongpu/param:
 - \$ ls -l include/picongpu/param
 - ⇒ The *.param files largely define the simulation setup





(4.1) Adjust setup parameters

- for now we will not go into details
- just to follow the workflow, we will adjust **one compile-time** parameter
- compile-time parameters are in include/picongpu/param
- let's see where the parameter files are:

```
cd include/picongpu/param
ls -1
```

let's edit the laser configuration vim incidentField.param

66 change a_0 from 8.0 to 5.0





(4.2) Adjust setup parameters

- iust to follow the workflow, we will adjust **one run-time** parameter
- run-time parameters are in etc/picongpu/*.cfg
- let's see where the parameter files are:

```
cd $HOME/picInputs/LWFA
cd etc/picongpu/
ls -l
```

let's edit the 4.cfg

```
vim 4.cfg
```

68 remove checkpointing and add mapped memory strategy

```
TBG_openPMD="--openPMD.period 100 \\
             --openPMD.file simData \\
             --openPMD.ext bp \\
             --openPMD.dataPreparationStrategy mappedMemory"
   · 2024-10-25
```







(5) compile setup

- go back to the root of your LWFA example cd \$HOME/picInputs/LWFA
- get compute resources for compiling: getDevice
- compile your setup pic-build
- better:
 pic-build -f 2> err.txt
 - Allows to read errors in err.txt (after compiling)
 - Using option -f ensures we do not use cached *.param files but the original ones during compile.
 Accidentally using cached files is a common source of errors.
 - -c ... fixes a current issue with CMake

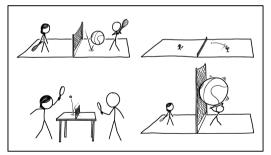






You can spend compile time to rethink your parameter choices (or check the latest xkcd...)

pic-build -f 2> err.txt



PARAMETERBALL IS A RAQUET GAME DIVIDED INTO FOUR QUARTERS, WITH BALL SIZE, COURT SIZE, AND NET HEIGHT RANDOMIZED EACH QUARTER.







We have a reservation

- in order to get your simulation to start, please use today's reservation
- the reservation is called:

ENCCS_day3

adjust or tbg template file

#SBATCH --exclusive

```
vim etc/picongpu/lumi-eurohpc/standard-g.tpl
replace line 25:
    #SBATCH --partition=!TBG_queue
with:
    #SBATCH --reservation=ENCCS_day3
```





- (6) Run setup
 - go back to LUMI head node do not run tbg from a compute node
 - submit job using tbg (assuming you changed the standard-g.tpl)
 tbg -s -t -c etc/picongpu/4.cfg \$SCRATCH/\$(whoami)/01_LWFA
 - tbg stand for Template Batch Generator and is our abstraction for setups for various clusters
 - it sets a submit command via ¬s (default on LUMI: sbatch)
 - it uses a template via -t (default on LUMI: standard-g.tpl)
 - it uses a setup configuration
 - does your job run?
 squeue -u \$USER
 scontrol show jobid <Your job's id>







Part III

Data analysis with openPMD and Jupyter

Analysing your PIConGPU simulation

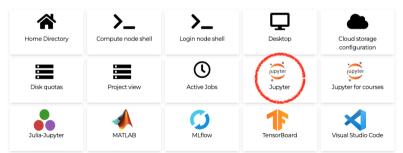
using jupyter, openPMD-api, ...

again use the LUMI web interface

www.lumi.csc.fi

- login with your credentials
- select Jupyter

Pinned Apps



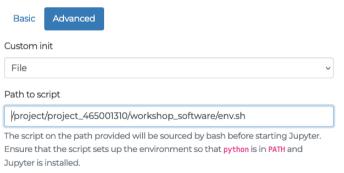






Analysing your PIConGPU simulation getting a jupyter job

- set reservation
- use 2 cores
- use advanced setting and use prepared python environment
- for now use your home directory as working directory



Launch



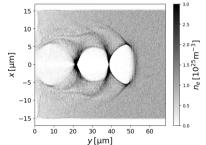




Analysing your PIConGPU simulation

run an example notebook

- copy notebook to your home directory
 cp /projappl/project_465001310/workshop_software/notebooks/LWFA_analysis.ipynb
 \$HOME/
- change the simulation directory to your simulation path



- execute it
- explore other data









Part IV

Adjusting simulation parameters

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simulation.param

- contains grid layout Δx , Δy , Δz , Δt
- reference density BASE_DENSITY_SI
- number of particles per cell to be initialized TYPICAL PARTICLES PER CELL





speciesInitialization.param

contains a pipeline that described how macro-particles are initialized

```
using InitPipeline = pmacc::mp_list<
    ...>;
```

particles can be created based on a density profile

```
CreateDensity<densityProfiles::Gaussian,
    startPosition::Random,
    PIC_Electrons>
```

or derived from an already existing particle distribution

```
Derive<PIC_Electrons, PIC_Ions>
```

particle attributes can be manipulated to set charge states, temperatures, drifts, ...







density.param

- LWFA contains only a Gaussian density profile
- let's have a look at it ...
- more density profiles are available, most flexible is:

```
struct FreeFormulaFunctor
    HDINLINE float X operator()(const floatD 64% position SI,
        const float3 64& cellSize SI) {
        float X = 1.0 \times -5.0 \times \text{math::abs(position SI.y()} - 0.0002 \times);
        s *= float X(s >= 0.0):
        return s;
};
using FreeFormula = FreeFormulaImpl<FreeFormulaFunctor>;
```



incidentField.param

- LWFA uses Gaussian laser profile
- **at** the very end:

```
using XMin = profiles::None;
using XMax = profiles::None;
using YMin = PARAM_LASERPROFILE;
using YMax = profiles::None;
using ZMin = profiles::None;
using ZMax = profiles::None;
```

laser definition:

```
using GaussianProfile = profiles::GaussianPulse<GaussianPulseParam>;
```

let's have a detailed look





Task

run your own mini-simulation campaign

- based on your initial LWFA setup, either:
- simulate half and double the density and study the influence on the wakefield
- lacksquare simulate $a_0=1.0$ and $a_0=0.2$ and see the effect on the wakefield
- play around with any other parameter you like







Part V

Time for your questions

- What is ...?
- How do I do ...?
- Is it possible to ...?
 Yes! (with enough time and effort ;-)
- Where do I get more information on ...? In order of precedence:
 - The manual: https://picongpu.readthedocs.io
 - Old issues https://github.com/ComputationalRadiationPhysics/picongpu/issues?q=is%3Aissue
 - Open a new issue







Part VI

Starting from another examples

1

Explore other default examples

setup your own example - we are her to help you

Kelvin Helmholtz instability

```
KelvinHelmholtz
use 4.cfg and add openPMD output
https://github.com/ComputationalRadiationPhysics/picongpu/tree/dev/share/
picongpu/examples/KelvinHelmholtz
```

Solid density foil target for proton acceleration

```
FoilLCT
use 4.cfg or 8.cfg
only uses 2D
https://github.com/ComputationalRadiationPhysics/picongpu/tree/dev/share/
picongpu/examples/FoilLCT
```



