



Vlaanderen
is supercomputing

JupyterHub Lunchbox session

Poll



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I intend to access JupyterHub ...

to develop CPU/GPU
software

to pre- or post-process
data interactively

to create
visualizations

for educational
purposes

I am ...

already using
Jupyter Notebooks
for my work

planning to use
Jupyter Notebooks
in the future

How frequently do you use Jupyter Notebooks?

Rarely

Few times
a week

Every day

General – Best practices

- Easy use of Jupyter Notebooks on the HPC facilities
 - Interactive session
- Best used for:
 - Data exploration
 - Visualization
 - Education
 - According to standard rules for HPC courses
 - Prototyping/development
- **Don't use with:**
 - Version control
 - Complicated dataflows

Getting started

- Open your browser and go to <https://jupyterhub.hpc.kuleuven.be/hub>
- Select the required resources
 - 1, 4 or 12h
 - CPU or GPU
 - Best lowest possible resources, increase if necessary
- Select desired project
- Standard location is \$VSC_DATA
- If server doesn't start in 5 mins, process will be killed → try again

Server Options

Select a job profile:

CPU - 1 core, 5 GB, 1 hour

Select a project:

lpt2_sysadmin (28136.67 credits available)

For further information on the service (status, bugs, announcements) please visit:
<https://jupyterhub-doc.readthedocs.io>

Projects queried and page generated at 2021-08-31 15:43:22

Start

Kernels

- Two kernels available
 - Python 3.6.8
 - R 3.6.3
 - RStudio also available in NX, but kernel is better for intense calculations
- Limited packages available, recommended to create your own kernel

Creating your own kernel

- R or Python kernel

- Use conda:

- Install Miniconda in \$VSC_DATA:

Python: https://docs.vscentrum.be/en/latest/software/python_package_management.html#install-miniconda

R: https://docs.vscentrum.be/en/latest/software/r_package_management.html#installing-miniconda

- Create new conda environment or use existing environment
 - Install necessary packages
\$ conda install <package_name>

Creating your own kernel: Python

- Activate your environment:

\$ source activate <envname>

- Include the IPython kernel, as well as ipython_genutils

\$ conda install ipykernel

\$ conda install ipython_genutils

- Install environment in *\$VSC_HOME/.local* with ipykernel:

\$ python -m ipykernel install --prefix=\${VSC_HOME}/.local/ --name '<envname>' --display-name '<name>'

- New packages can be added from within JupyterHub.

\$ conda install <package_name>

- only one conda command per cell
- Restarting the kernel is necessary

Creating your own kernel: R

- Activate your environment:

\$ source activate <envname>

- Install IRKernel and jupyter_client:

\$ conda install jupyter_client r-irkernel -c conda-forge

- Install the environment to `$VSC_HOME/.local` with IRkernel:

Rscript -e 'IRkernel::installspec(prefix="\$VSC_HOME/.local/", name=" <envname>", displayname=" <display_name>")'

- Conda packages cannot be installed from within Jupyter notebook

Removing a kernel

- Remove kernel in `$VSC_HOME/.local`
`$ rm $VSC_HOME/.local/share/jupyter/kernels/<kernel_name> -r`
- Be careful: removing a conda env and not removing the kernel as above will result in a non-working kernel. No error/warning is given in the Jupyter Notebook

Limitations

- Not possible to load any extra modules.
 - Have a look for a conda/pip alternative if possible.
 - In case you have a specific case where you can only use a specific VSC module, please contact the servicedesk
- Other executables (e.g. Tensorboard) cannot be used in Jupyter at the moment

New kernel installation and notebook examples

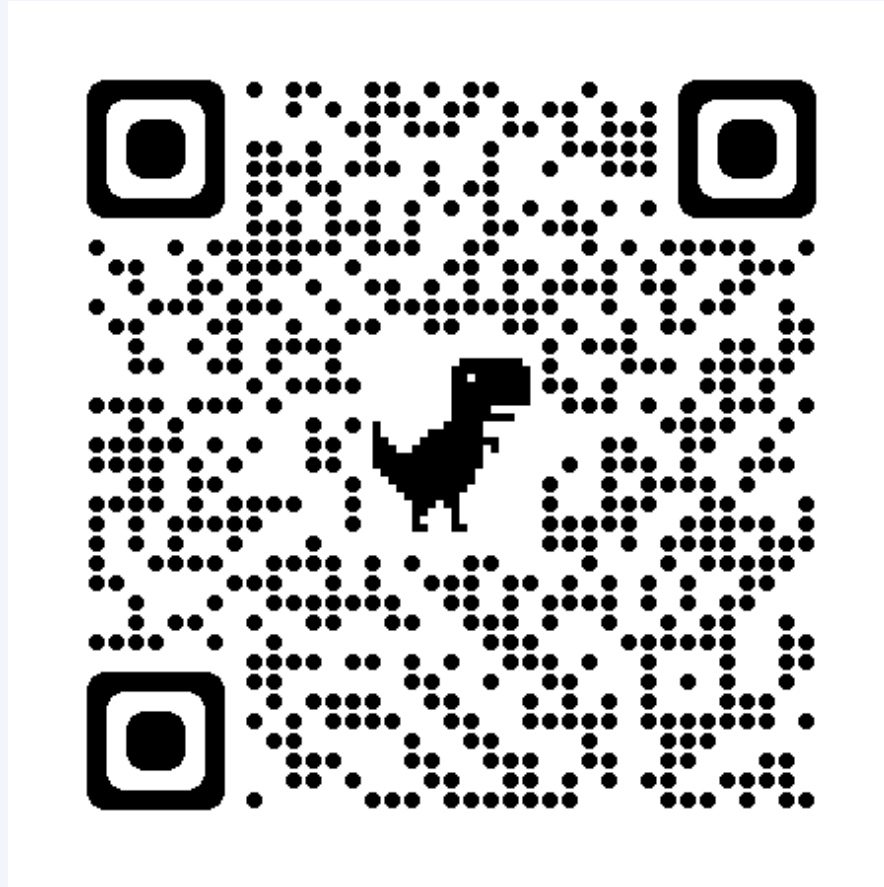
Python

- MNIST on GPU

R

- K-means clustering

Poll



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I intend to use ...

resources with
CPUs only

resources with
GPUs

Desired number of cores when using CPU resources:

Not applicable

1

2

4

More

Useful duration of the JupyterHub session

1 hour

4 hours

12 hours

Which kernel do you use?

Python

R

Bash