



(virtual) HPC Champions meeting (UK)

Tue Dec 15th 2020

EESSI

EUROPEAN ENVIRONMENT FOR
SCIENTIFIC SOFTWARE INSTALLATIONS

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whoami

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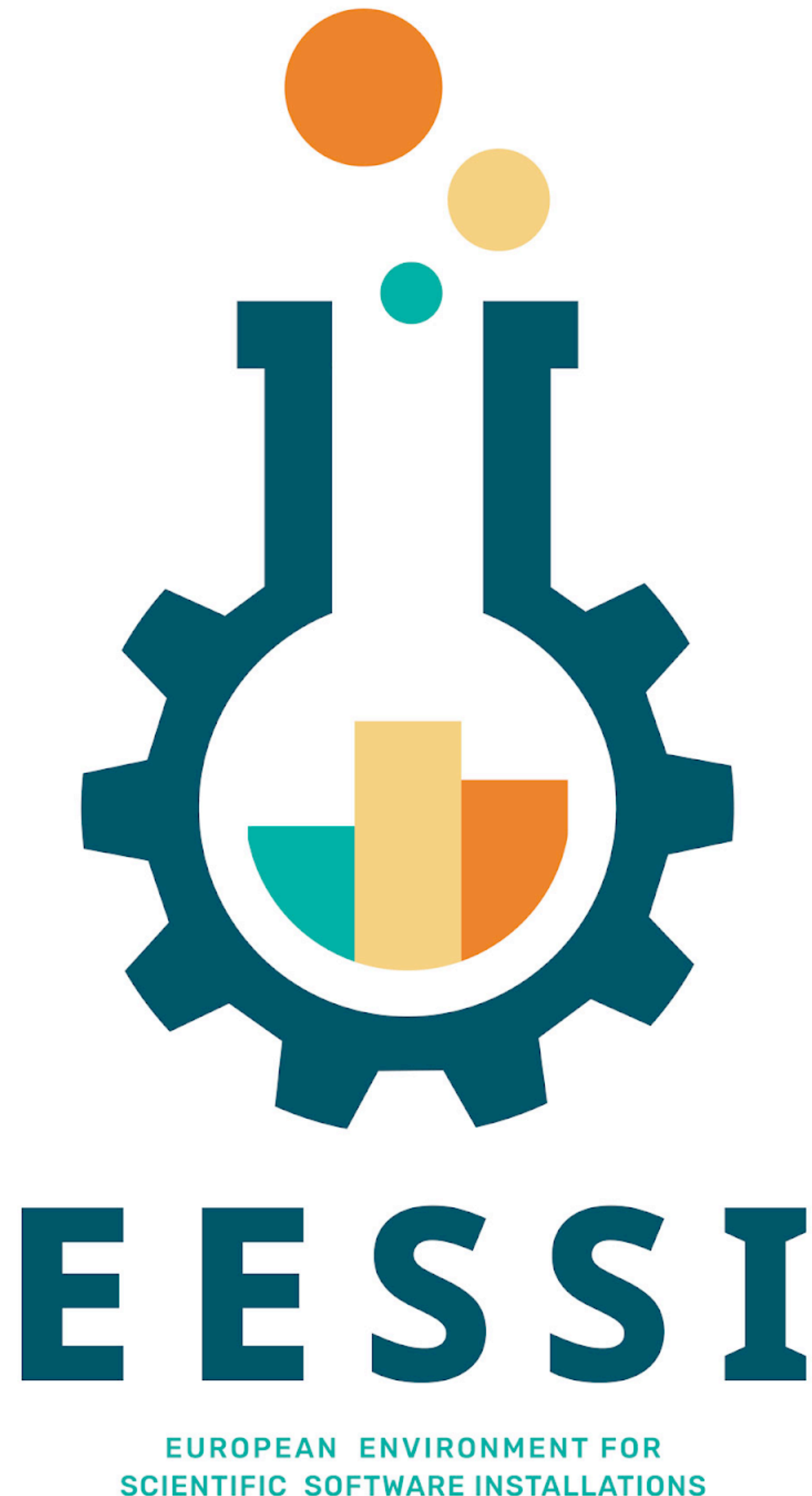
@kehoste



@boegel



- Computer scientist from Belgium
- Masters & PhD from Ghent University
- **HPC system administrator at UGent since 2010**
- User support & training, **software installations** (RSE?)
- Big fan of FOSS, family, loud music, (good) beer, dad jokes, stickers, ...
- Doesn't like C++, CMake, SCons, Bazel, TensorFlow, OpenFOAM, ...



- **Who** is involved in EESSI?
- **What** is the project about?
- **Why** did we start it?
- **How** are we tackling the problem?
- What are the **challenges**?
- Which **FOSS projects** do we use?
- What is the **current status**?
- **Live demo(s)!**

EESSI in a nutshell



- **European Environment for Scientific Software Installations**
(EESSI, pronounced as "easy")
- Collaboration between different partners in HPC community
- Goal: **building a common performant scientific software stack**
for HPC systems & beyond (cloud, workstations, ...)
- Grass roots project

<https://eessi-hpc.org>

<https://github.com/EESSI>

<https://eessi.github.io/docs/pilot>

 *[@eessi_hpc](https://twitter.com/eessi_hpc)*

Project partners



UNIVERSITY OF TWENTE.



UiO : **University of Oslo**



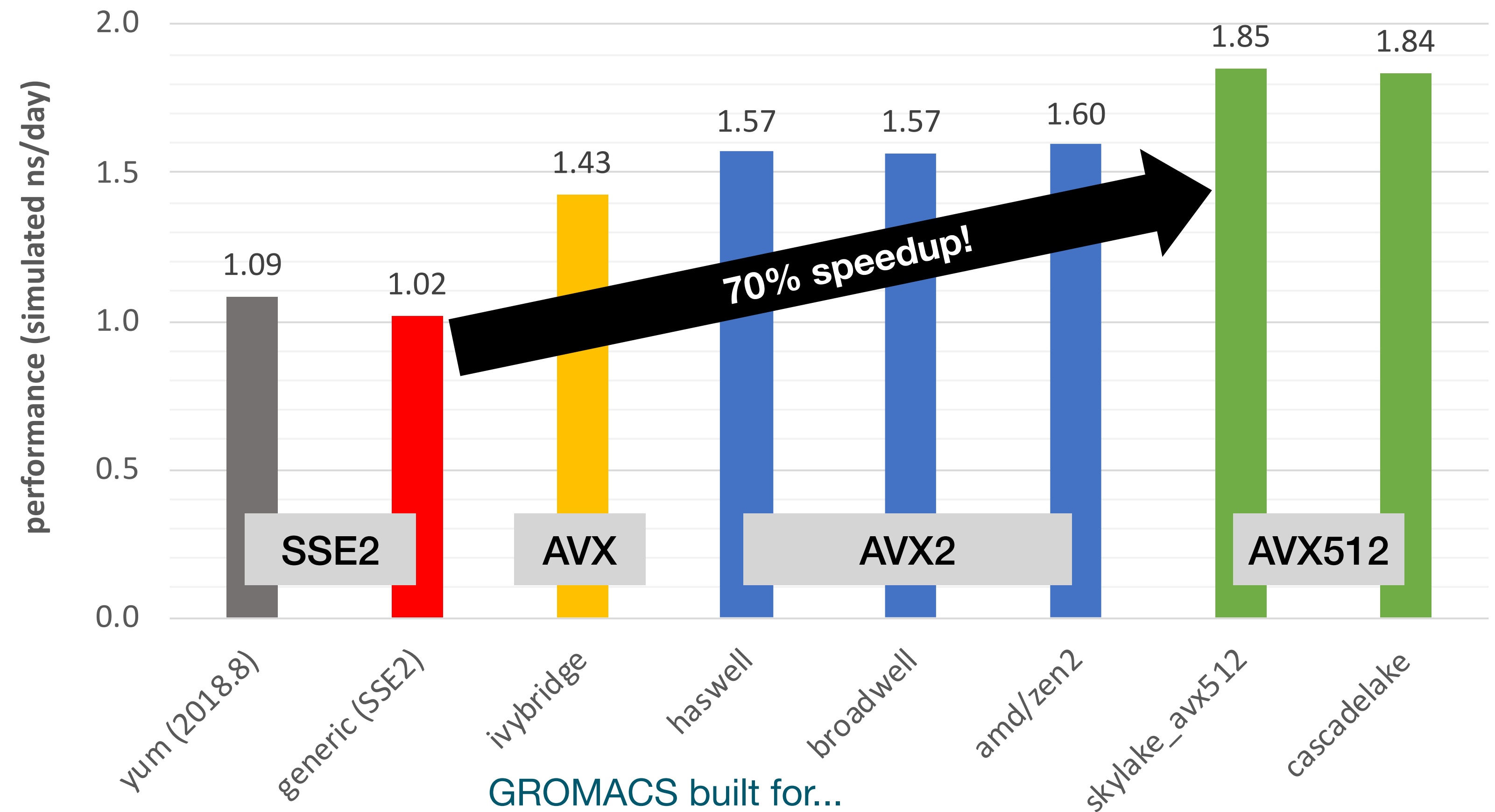
There's a storm coming... or are we in it already?

- Explosion of open source scientific software in recent years
- Increasing variety in CPUs: Intel, AMD, Arm, POWER, RISC-V
- Various types of accelerators: NVIDIA & AMD GPUs, Intel Xe, ...
- Rise of the cloud: Amazon EC2, Microsoft Azure, Google, Oracle, ...
- In stark contrast: available manpower in HPC support teams...

Keeping the P in HPC

- **Software should be optimised for the system it will run on**
- Impact on performance is often significant for scientific software

- Example: GROMACS 2020.1
(PRACE benchmark, Test Case B)
- Metric: (simulated) ns/day,
higher is better
- Test system: dual-socket
Intel Xeon Gold 6420
(Cascade Lake, 2x18 cores)



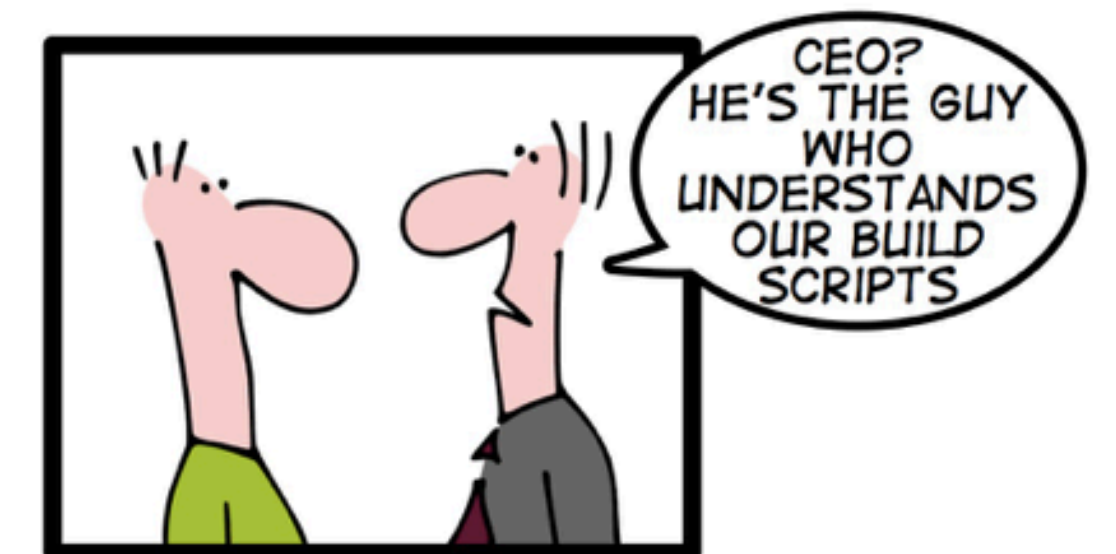
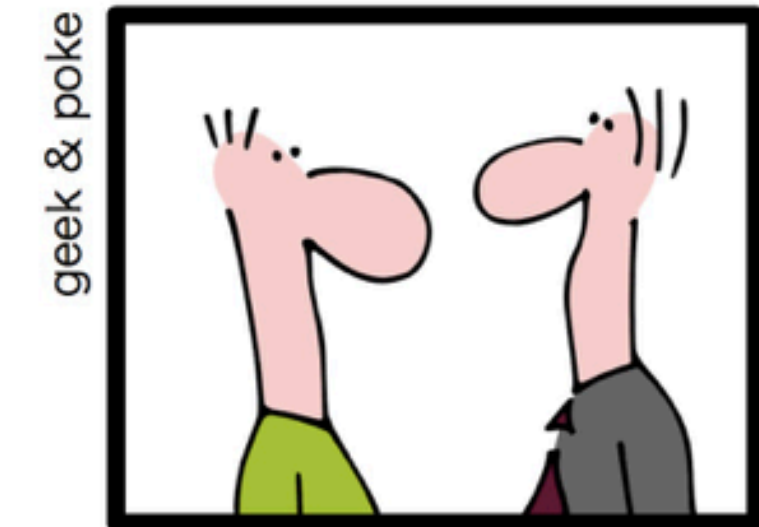
Getting Scientific Software Installed

```
INSTALL.SH  
#!/bin/bash  
  
pip install "$1" &  
easy_install "$1" &  
brew install "$1" &  
npm install "$1" &  
yum install "$1" & dnf install "$1" &  
docker run "$1" &  
pkg install "$1" &  
apt-get install "$1" &  
sudo apt-get install "$1" &  
steamcmd +app_update "$1" validate &  
git clone https://github.com/"$1"/"$1" &  
cd "$1";./configure;make;make install &  
curl "$1" | bash &
```

<https://xkcd.com/303>

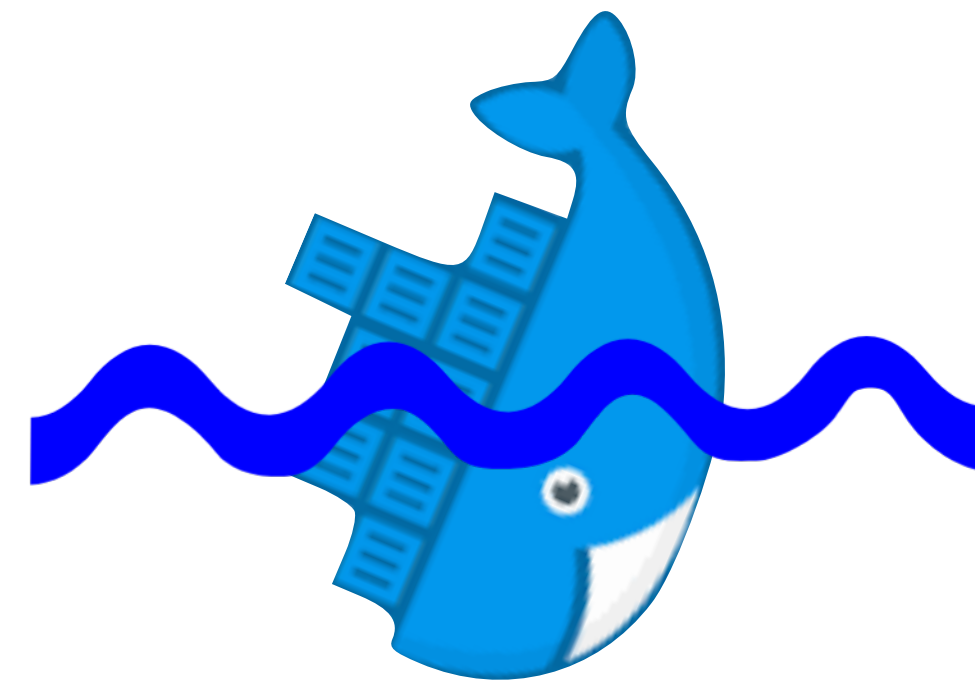


<https://xkcd.com/1654>



HOW TO BECOME INVALUABLE

What about... containers?



- Containers are (still) problematic in an HPC context
 - Integration with system resources & services is not a solved problem (MPI, GPUs, ...)
- "Native performance" really means "very little overhead"
 - That's very good, and a *requirement* for HPC, but not sufficient...
- Someone (you trust) has to build (and maintain) the container images you need!
 - Including the software you want to use, for your CPU architecture (x86_64, aarch64, ...)
 - Ideally, the image is properly optimised for the *specific* hardware you'll be using it on (performance vs "mobility of compute")
- **Containers are a symptom, not a cure...**



Scope & goals



- **Shared repository of scientific software installations**
- Collaborate, avoid duplicate work across HPC sites
- **Uniform way of providing software to researchers**
- Offer broad platform support (Linux, macOS, Windows via WSL)
- **Targets: laptops, personal workstations, HPC clusters, and the cloud**
- Support for different CPUs, interconnects, GPUs, etc.
- Focus on **performance**, automation, testing, tuning



<http://www.thecomicstrips.com/subject/The-Collaboration-Comic-Strips.php>

Pick a live demo! (software)



1



GROMACS

version 2020.1

Molecular dynamics simulation

PRACE benchmark (MPI)

2



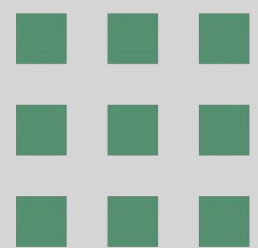
TensorFlow

version 2.3.1

Machine learning

Beginner tutorial with MNIST (single node)

3



OpenFOAM

version 8 (openfoam.org)

Computational Fluid Dynamics (CFD) simulation

motorBike tutorial case (MPI)

4



Bioconductor

version 3.11 (on top of R 4.0)

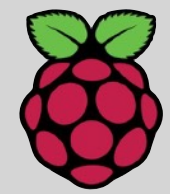
bioinformatics

Playing with DNA data (single node)

Pick a live demo! (hardware)



A



Raspberry Pi cluster

4x model 3B+



quad-core **Arm** Cortex-A53, 1GB RAM

64-bit Raspberry Pi OS

native CernVM-FS (built from source)

B



Amazon EC2

heterogeneous Slurm cluster

created via Cluster-in-the-Cloud

Intel Xeon, AMD Rome, Graviton2

CentOS 8.3 , native CernVM-FS



Standard_D8s_v3 instance

8-core Intel Xeon (**different CPU generations**)

Ubuntu 18.04

C

native CernVM-FS, **installed from scratch**

Isambard2 cluster (XCI partition)

Cray XC50 @ GW4 / UK Met Office

Marvell Thunder X2 Arm64 CPUs (> 20k cores)

Cray Linux Environment 7 (SLES 15)

CernVM-FS via **Singularity** container

D

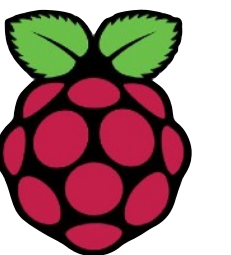


Spoiler demo



GROMACS

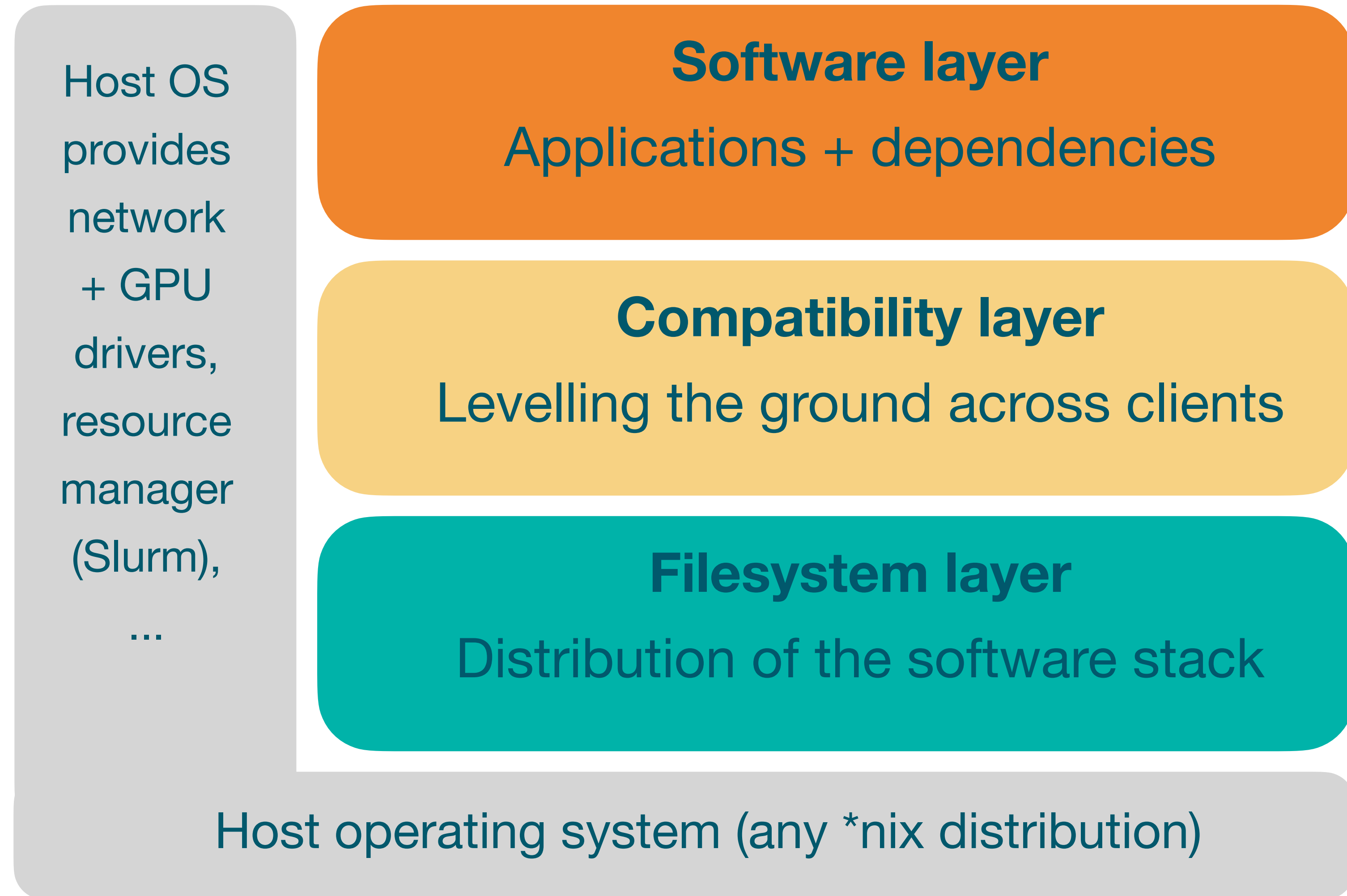
*Let's run a molecular
dynamics simulation on
a Raspberry Pi cluster!*



E E S S I

EUROPEAN ENVIRONMENT FOR
SCIENTIFIC SOFTWARE INSTALLATIONS

High-level overview of EESSI project

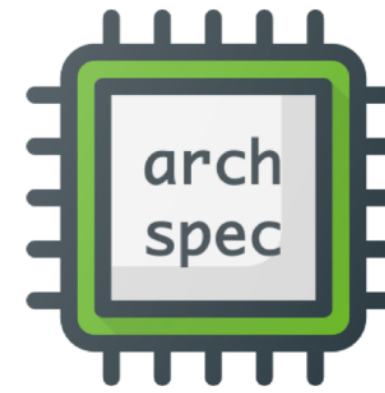


EESSI is powered by FOSS



- Installation tool for scientific software
- Optimises for build host (by default)
- Supports over 2,000 software pkgs

<https://easybuilders.github.io/easybuild>



- Python library
- Detect processor type
- Check compatibility with host CPU

<https://github.com/archspec>



- Environment modules tool (written in Lua)
- Intuitive access to software installations
- Multiple software versions side-by-side

<https://lmod.readthedocs.io>



<https://wiki.gentoo.org/wiki/Project:Prefix>

- Linux distribution, installs from source
- Prefix project: install packages in <prefix>
- Supports x86_64, Arm64, POWER, ... & Linux, macOS



CernVM-FS

- Software distribution service
- Scalable, read-only, globally distributed filesystem
- Mount filesystem over HTTP

<https://cernvm.cern.ch/fs>



- Regression testing framework for HPC
- Verify correctness
- Check performance
- Tests are implemented in Python

<https://reframe-hpc.rtd.io>



- Automation
- Configuration mgmt



Singularity

- Build isolation
- Easy access for clients



HashiCorp

Terraform

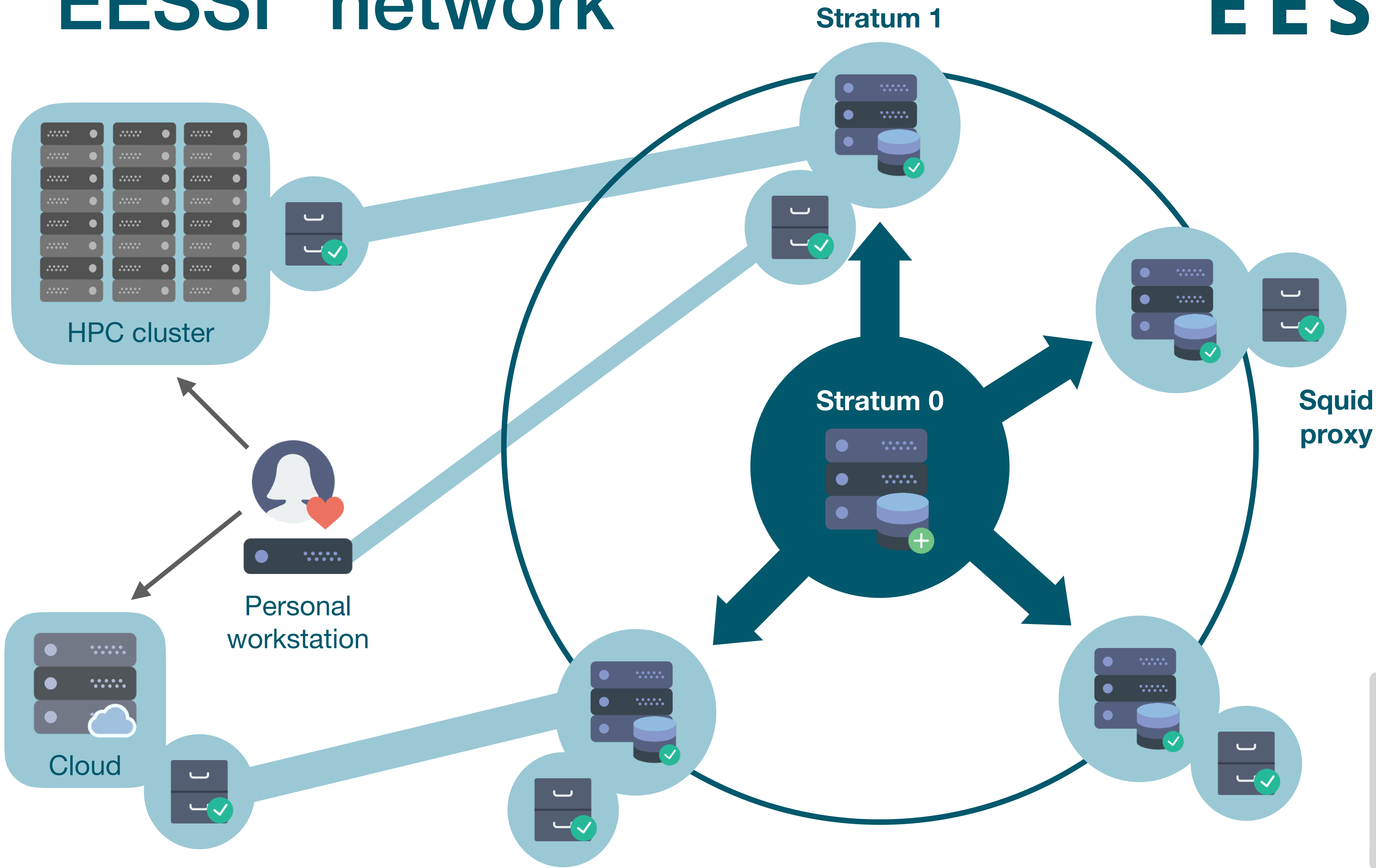
- Creating cloud instances on demand

Inspiration for EESSI



- EESSI concept is **heavily** inspired by Compute Canada software stack
- Shared across 5 major national systems in Canada + a bunch of smaller ones
- 3 layers: CernVM-FS / ~~Nix~~ Gentoo Prefix / EasyBuild + Lmod
- See paper by Maxime Boissonneault & co at PEARC'19 (PDF available [here](#))
“Providing a Unified Software Environment for Canada’s National Advanced Computing Centers”
- See also Maxime’s talk at 5th EasyBuild User Meeting ([slides](#) - [recorded talk](#))
and the Compute Canada [documentation](#)

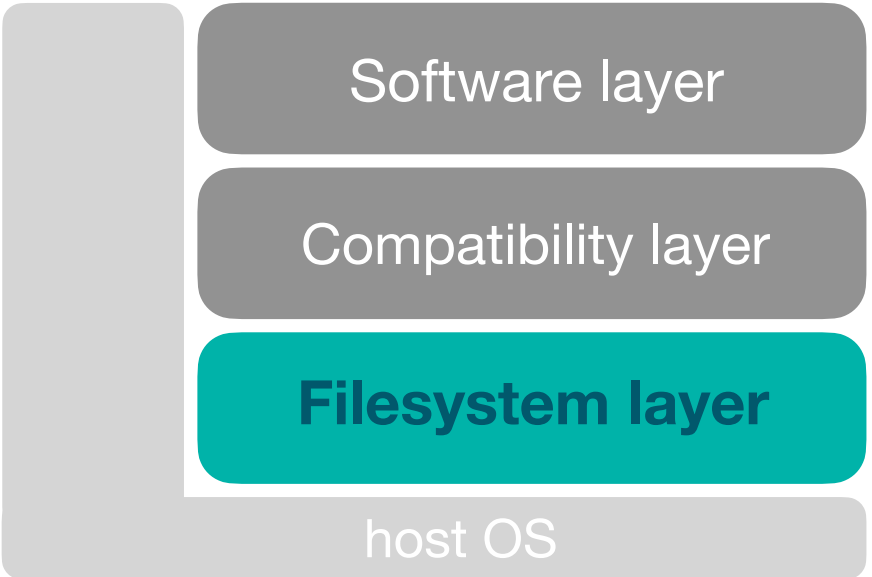
EESSI "network"



powered by



CernVM-FS



EESSI "network"



Key messages:

- CernVM-FS provides a scalable, reliable setup
- Distributed access via HTTP (so firewall-friendly)
- **Same software stack available everywhere!**

powered by



CernVM-FS

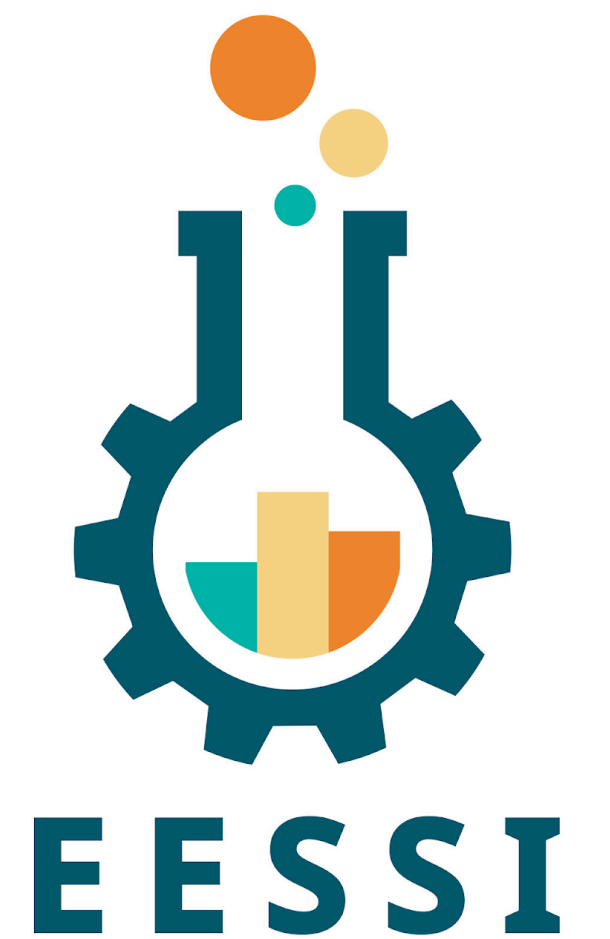


Compatibility layer

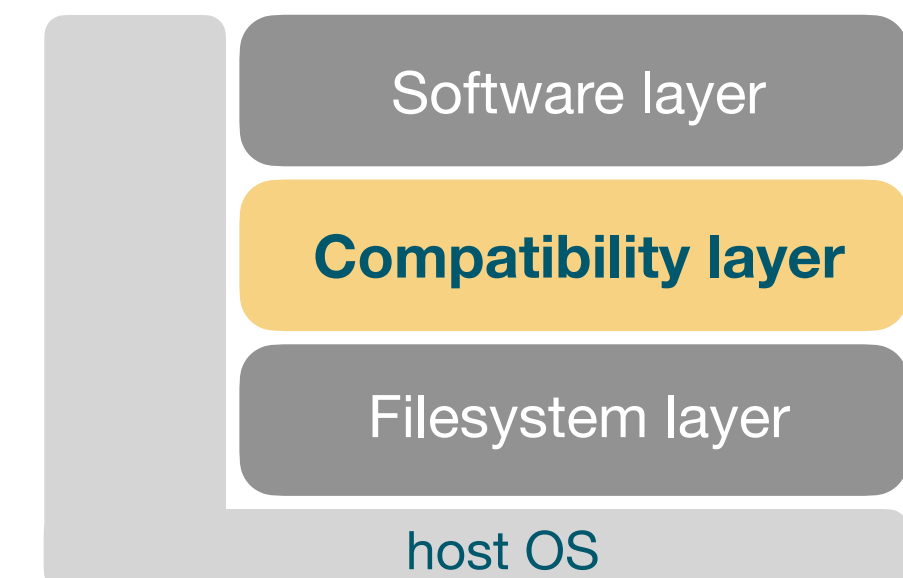
- Set of tools & libraries installed in non-standard location
- Using Gentoo's package manager Portage
- Limited to low-level stuff, incl. glibc
- Only targets a supported processor **family** (x86_64, Arm64)
- **Levels the ground for different client operating systems** (Linux distros, macOS)
- Currently in pilot repository:

`/cvmfs/pilot.eessi-hpc/2020.12/compat/linux/aarch64`

`/cvmfs/pilot.eessi-hpc/2020.12/compat/linux/x86_64`

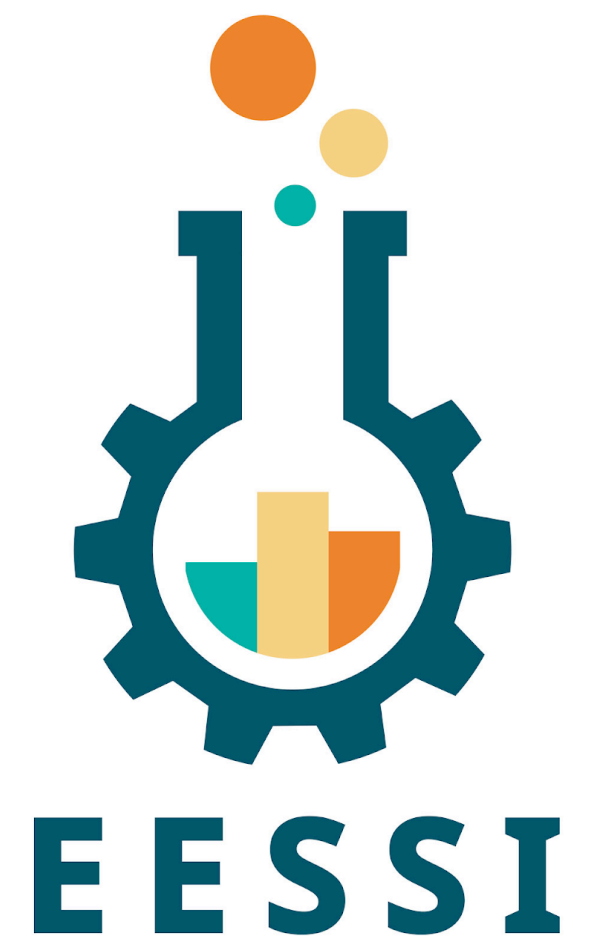


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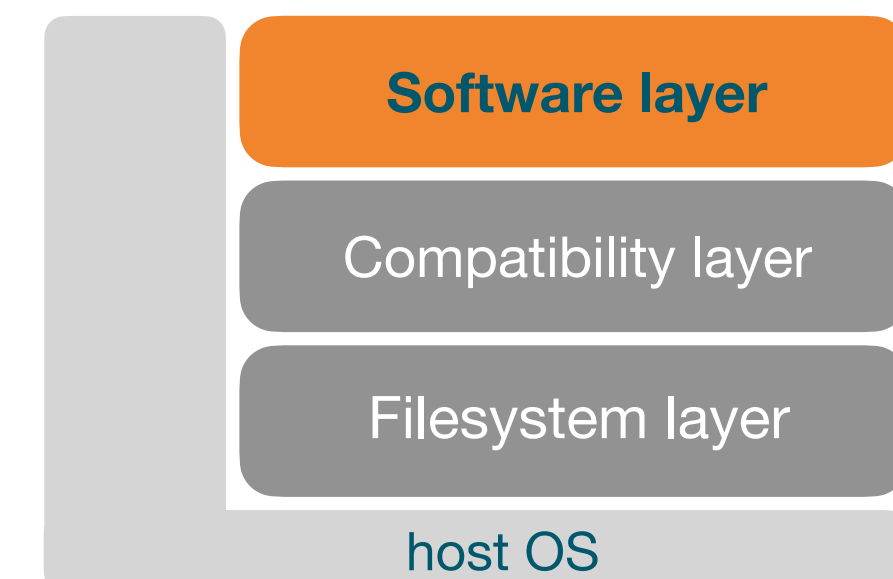
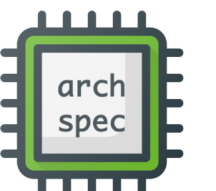


Software layer

- Provides scientific software applications, libraries, and dependencies
- **Optimised for specific CPU microarchitectures** (Intel Haswell, ...)
- **Leverages libraries from compatibility layer** (not from host OS)
- Installed with EasyBuild, incl. environment module files
- Lmod environment modules tool is used to access installations
- Different subdirectories: one per target CPU microarchitecture
- **Best subdirectory for host is picked automatically** via archspec



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Current status: pilot repository



- Ansible playbooks, scripts, docs at <https://github.com/eessi>
- Stratum 0 @ Univ. of Groningen + two Stratum 1 servers
- Compatibility layer for both x86_64 and aarch64 (only Linux clients, for now)
- Software (CPU-only): Bioconductor, GROMACS, OpenFOAM, TensorFlow
- Hardware targets:
 - x86_64/generic, intel/haswell, intel/skylake_avx512, amd/zen2
 - aarch64/generic, aarch64/graviton2, **aarch64/thunderx2**

**NOT FOR
PRODUCTION USE!**

Try it yourself: <https://eessi.github.io/docs/pilot>

From zero to science in 3 steps



Step 1: Access the (pilot) EESSI CernVM-FS repository

Option 1: Native CernVM-FS installation (requires admin privileges)

- Install CernVM-FS, see <https://cernvm.cern.ch/fs>
- Also install configuration files for EESSI repositories, available at <https://github.com/EESSI/filesystem-layer/releases>
- Run "cvmfs_config setup"

Option 2: Use a Singularity container (no admin rights needed!)

- CernVM-FS repositories can be mounted via Singularity's `--fusemount` option
- See detailed walkthrough at <https://eessi.github.io/docs/pilot>

From zero to science in 3 steps

Step 1: Access the (pilot) EESSI CernVM-FS repository

Option 1 (example): native CernVM-FS installation on fresh (x86_64) RHEL 8.2 system



```
# install CernVM-FS (see https://cernvm.cern.ch/fs/)
sudo yum install -y https://ecsft.cern.ch/dist/cvmfs/cvmfs-release/cvmfs-release-latest.noarch.rpm
sudo yum install -y cvmfs

# install CernVM-FS configuration files for EESSI repositories (see https://github.com/EESSI/filesystem-layer)
wget https://github.com/EESSI/filesystem-layer/releases/download/v0.2.3/cvmfs-config-eessi-0.2.3-1.noarch.rpm
sudo yum install -y cvmfs-config-eessi-0.2.3-1.noarch.rpm

# create local CernVM-FS configuration file (direct access, no proxy; 10GB for CernVM-FS cache)
sudo bash -c "echo 'CVMFS_HTTP_PROXY=DIRECT' > /etc/cvmfs/default.local"
sudo bash -c "echo 'CVMFS_QUOTA_LIMIT=10000' >> /etc/cvmfs/default.local"

# set up CernVM-FS
sudo cvmfs_config setup

# access EESSI pilot repository
ls /cvmfs/pilot.eessi-hpc.org/2020.10
```

From zero to science in 3 steps

Step 1: Access the (pilot) EESSI CernVM-FS repository

Option 2 (example): use Singularity to run Docker container to access EESSI



```
# configure Singularity (bind mounts + home directory)
mkdir -p /tmp/$USER/{var-lib-cvmfs,var-run-cvmfs,home}
export SINGULARITY_BIND="/tmp/$USER/var-run-cvmfs:/var/run/cvmfs,/tmp/$USER/var-lib-cvmfs:/var/lib/cvmfs"
export SINGULARITY_HOME="/tmp/$USER/home:/home/$USER"

# values to pass to --fusemount (EESSI config + pilot repositories)
export EESSI_CONFIG="container:cvmfs2 cvmfs-config.eessi-hpc.org /cvmfs/cvmfs-config.eessi-hpc.org"
export EESSI_PILOT="container:cvmfs2 pilot.eessi-hpc.org /cvmfs/pilot.eessi-hpc.org"

# minimal Docker container from Docker Hub (includes CernVM-FS + EESSI configuration files)
export DOCKER_IMAGE="docker://eessi/client-pilot:centos7-$(uname -m)-2020.10"

# start shell in Singularity container (ignore the scary looking 'setxattr' warnings, they're harmless)
singularity shell --fusemount "$EESSI_CONFIG" --fusemount "$EESSI_PILOT" $DOCKER_IMAGE

# access EESSI pilot repository
ls /cvmfs/pilot.eessi-hpc.org/2020.10
```

From zero to science in 3 steps



Step 2: Set up your environment by sourcing the EESSI `init` script

```
# source the EESSI init script to set up your environment
$ source /cvmfs/pilot.eessi-hpc.org/2020.10/init/bash
Found EESSI pilot repo @ /cvmfs/pilot.eessi-hpc.org/2020.10!
Using x86_64/intel/haswell as software subdirectory.
Initializing Lmod...
Prepending /cvmfs/pilot.eessi-hpc.org/2020.10/software/x86_64/intel/haswell/modules/all to $MODULEPATH...

Environment set up to use EESSI pilot software stack, have fun!

[EESSI pilot 2020.10] $ echo $EESSI_PREFIX
/cvmfs/pilot.eessi-hpc.org/2020.10

[EESSI pilot 2020.10] $ echo $EESSI_SOFTWARE_SUBDIR
x86_64/intel/haswell
```


From zero to science in 3 steps



Step 3: Load the modules for the software you want to use, and go!

```
# check which modules are available
[EESSI pilot 2020.10] $ module avail gromacs

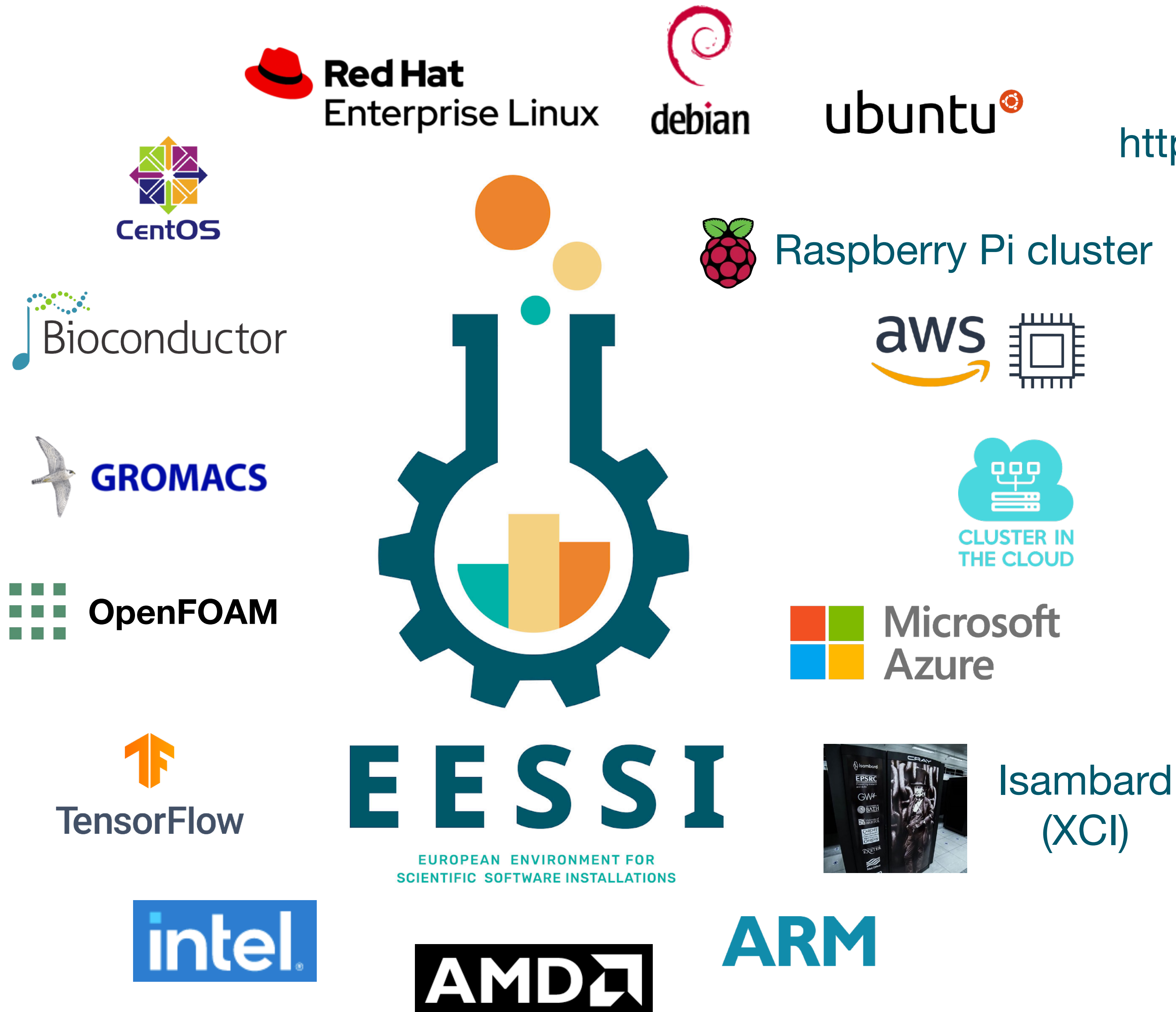
----- /cvmfs/pilot.eessi-hpc.org/2020.10/software/x86_64/intel/haswell/modules/all -----
GROMACS/2020.1-foss-2020a-Python-3.8.2

# load the module(s) for the software you want to use
[EESSI pilot 2020.10] $ module load GROMACS

# ready to compute!
[EESSI pilot 2020.10] $ gmx mdrun -s ion_channel.tpr -maxh 0.50 -rethway -noconfout -nsteps 1000
```

Demo time!

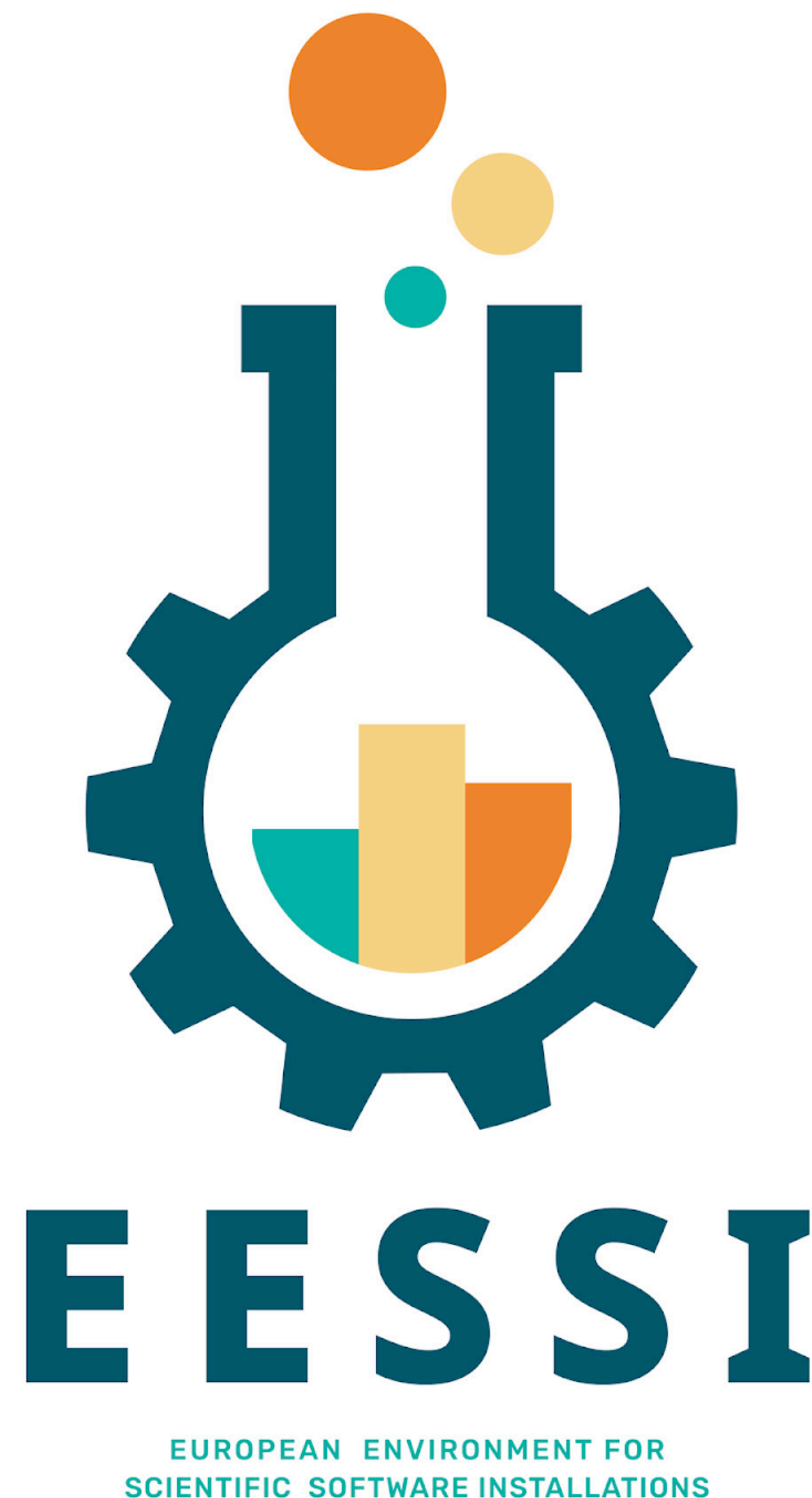
<https://github.com/EESSI/eessi-demo>



Future work



- Further improve pilot EESSI repository (~ monthly revisions)
- Identify problems, and fix them...
- **Automate** deployment of different EESSI layers (Ansible, Terraform, ...)
- **Testing** (with ReFrame) + continuous integration in GitHub Actions
- Let developers of scientific software validate the installation of *their* software
- Also support macOS / POWER / GPUs, add more software
- Solicit more manpower, get project funded to make it sustainable
- Work towards production setup...



Kenneth Hoste

kenneth.hoste@ugent.be

[@kehoste](#) (Twitter)

Website: <https://www.eessi-hpc.org>

Join our mailing list & Slack channel

<https://www.eessi-hpc.org/join>

Documentation: <https://eessi.github.io/docs>

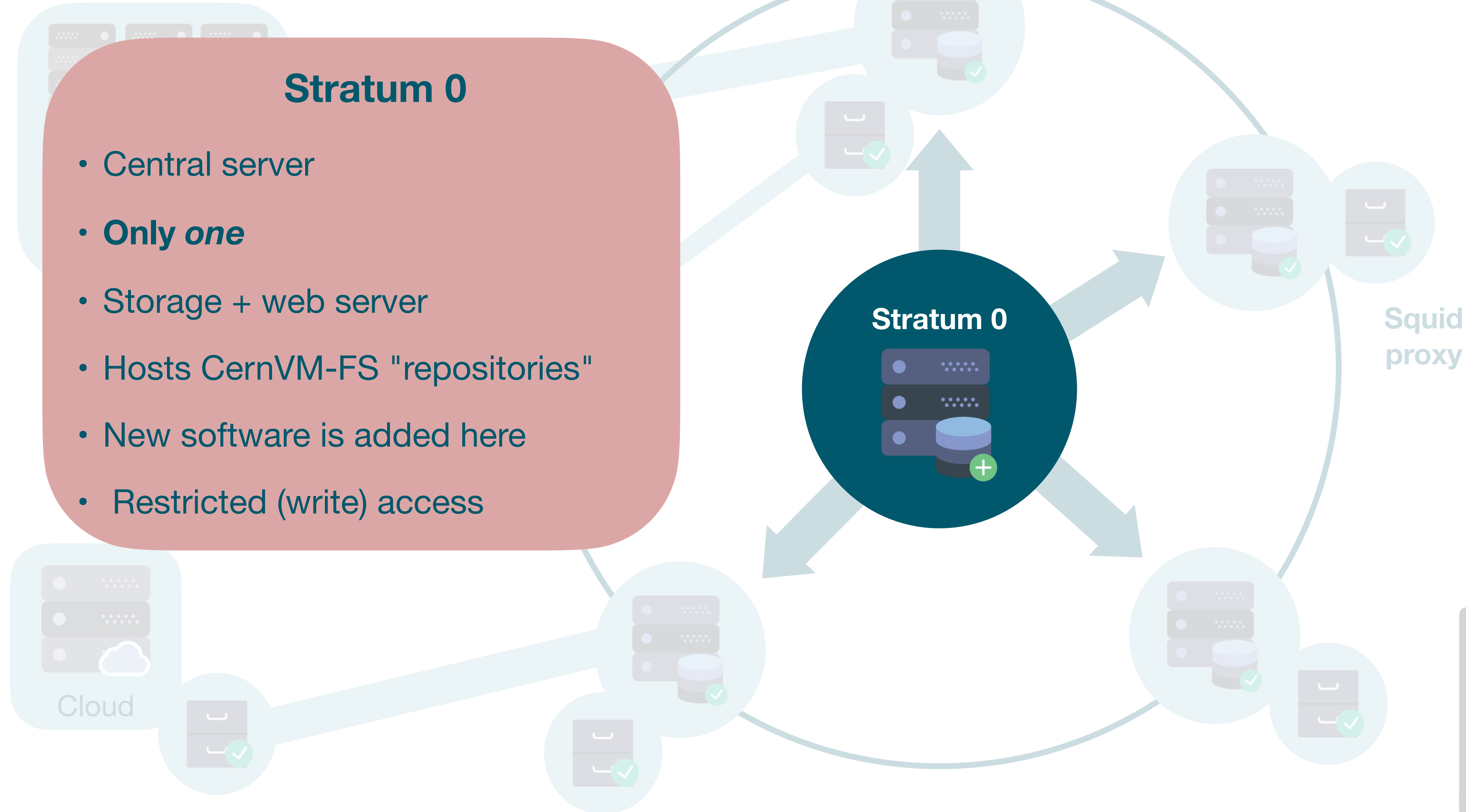
GitHub: <https://github.com/eessi>

Twitter: [@eessi_hpc](#)

Monthly online meetings (first Thursday, 2pm CET)

(BACKUP SLIDES)

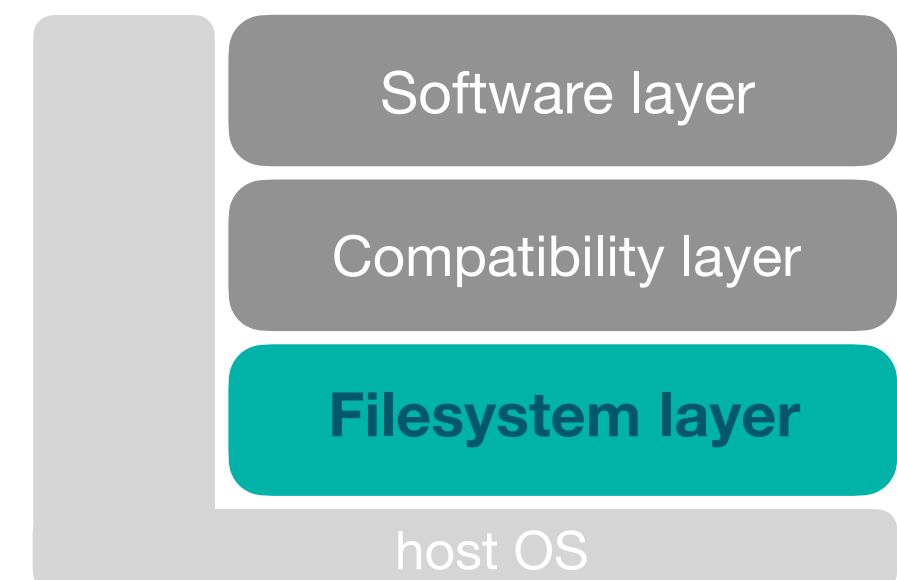
EESSI "network"



powered by



CernVM-FS



EESSI "network"

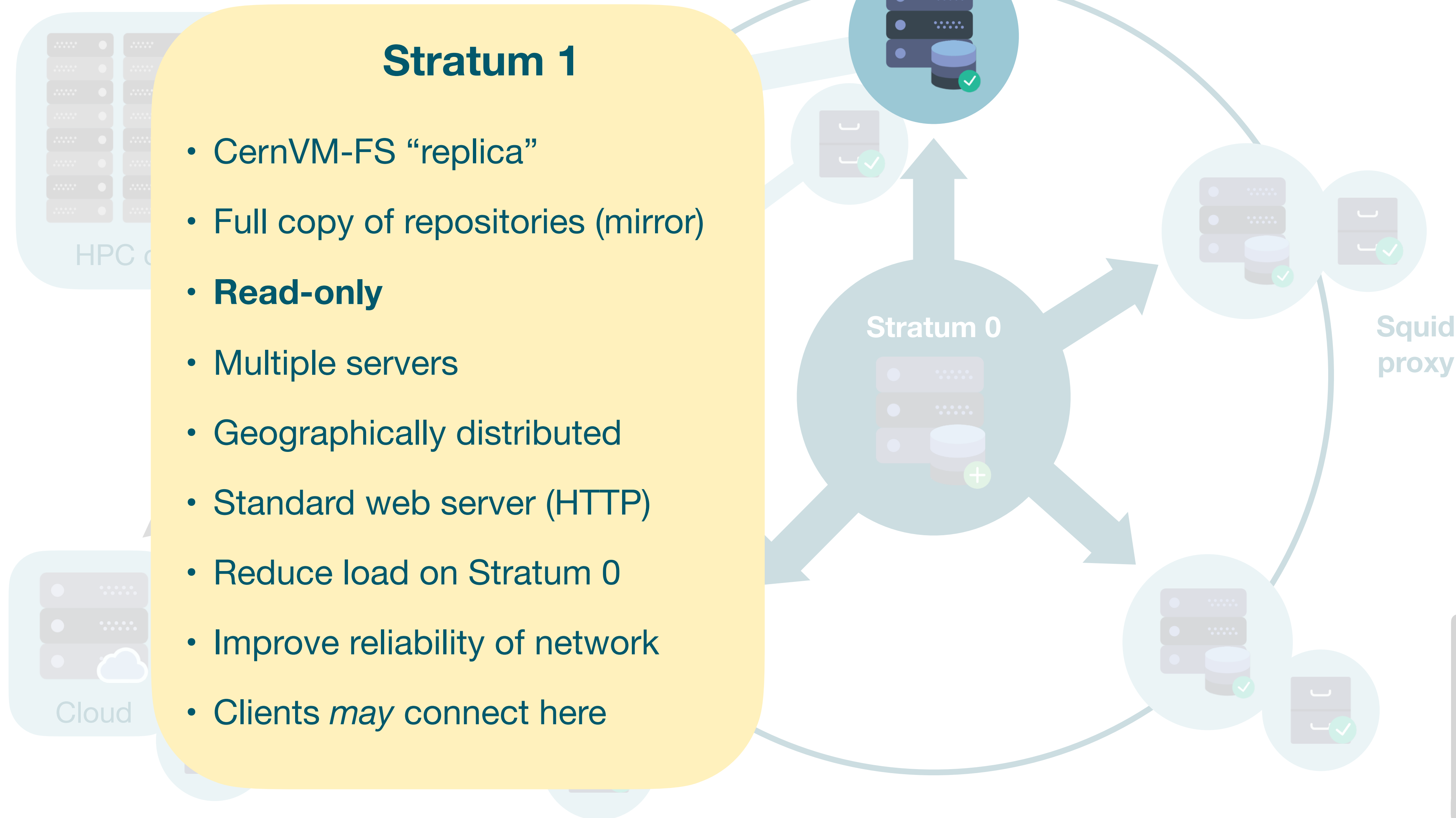


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CernVM-FS

- ## Stratum 1
- CernVM-FS “replica”
 - Full copy of repositories (mirror)
 - **Read-only**
 - Multiple servers
 - Geographically distributed
 - Standard web server (HTTP)
 - Reduce load on Stratum 0
 - Improve reliability of network
 - Clients *may* connect here



Squid proxy

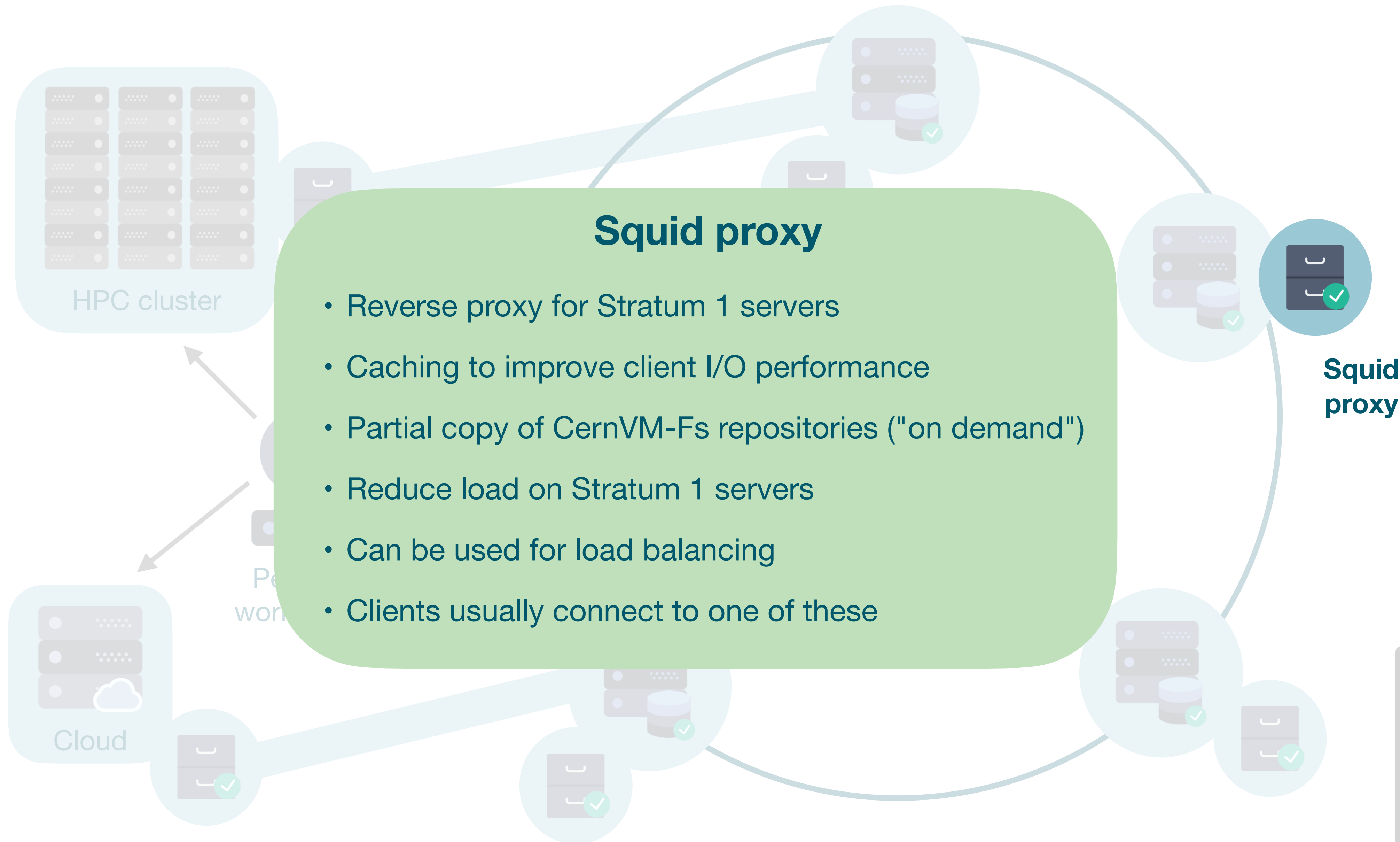
Software layer

Compatibility layer

Filesystem layer

host OS

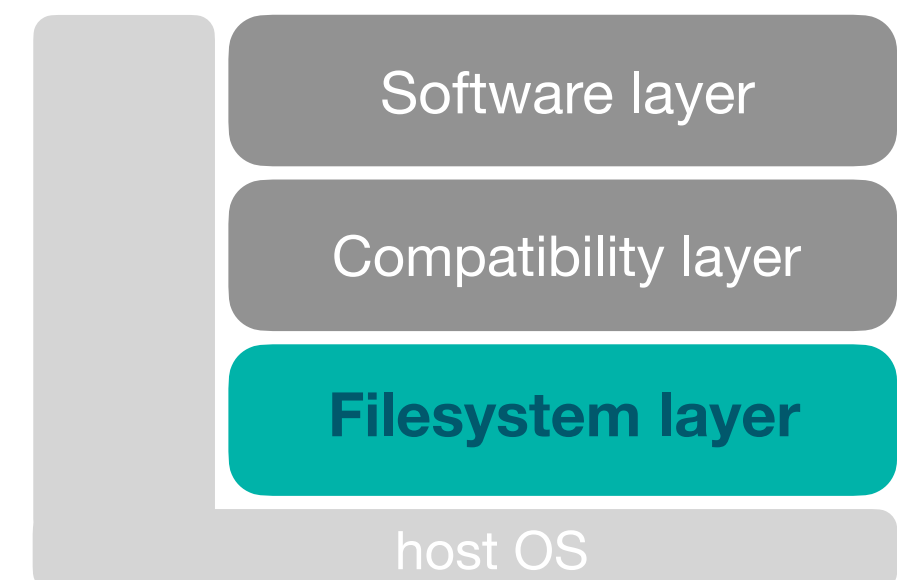
EESSI "network"



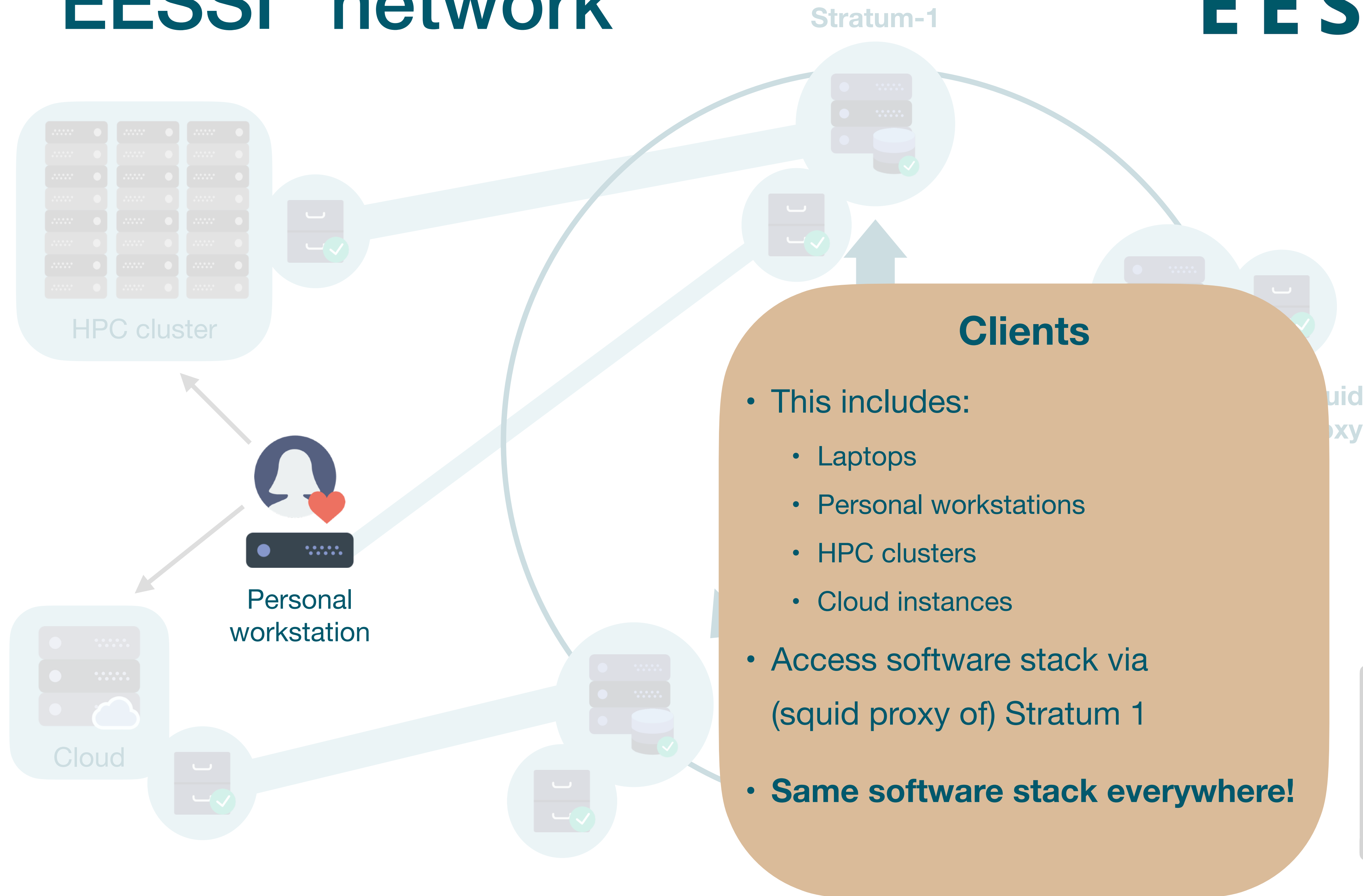
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CernVM-FS



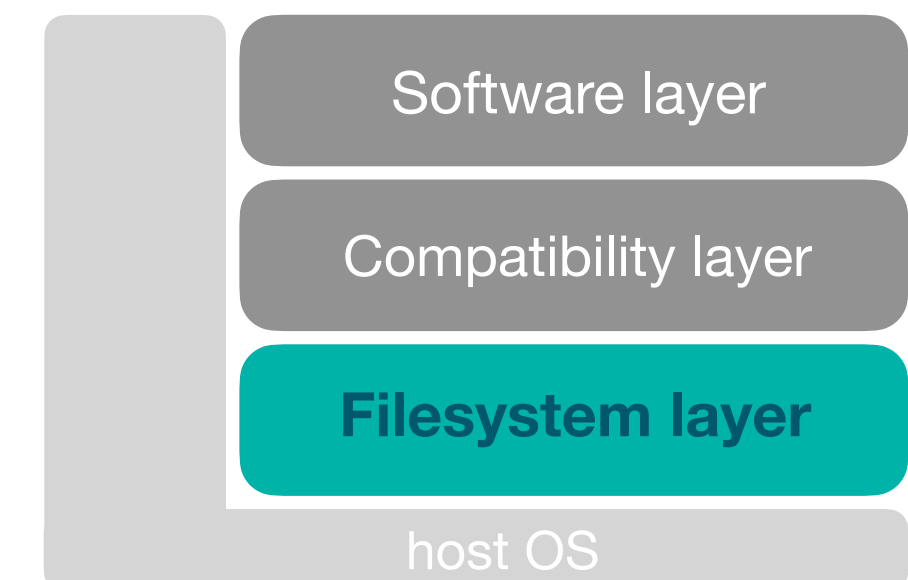
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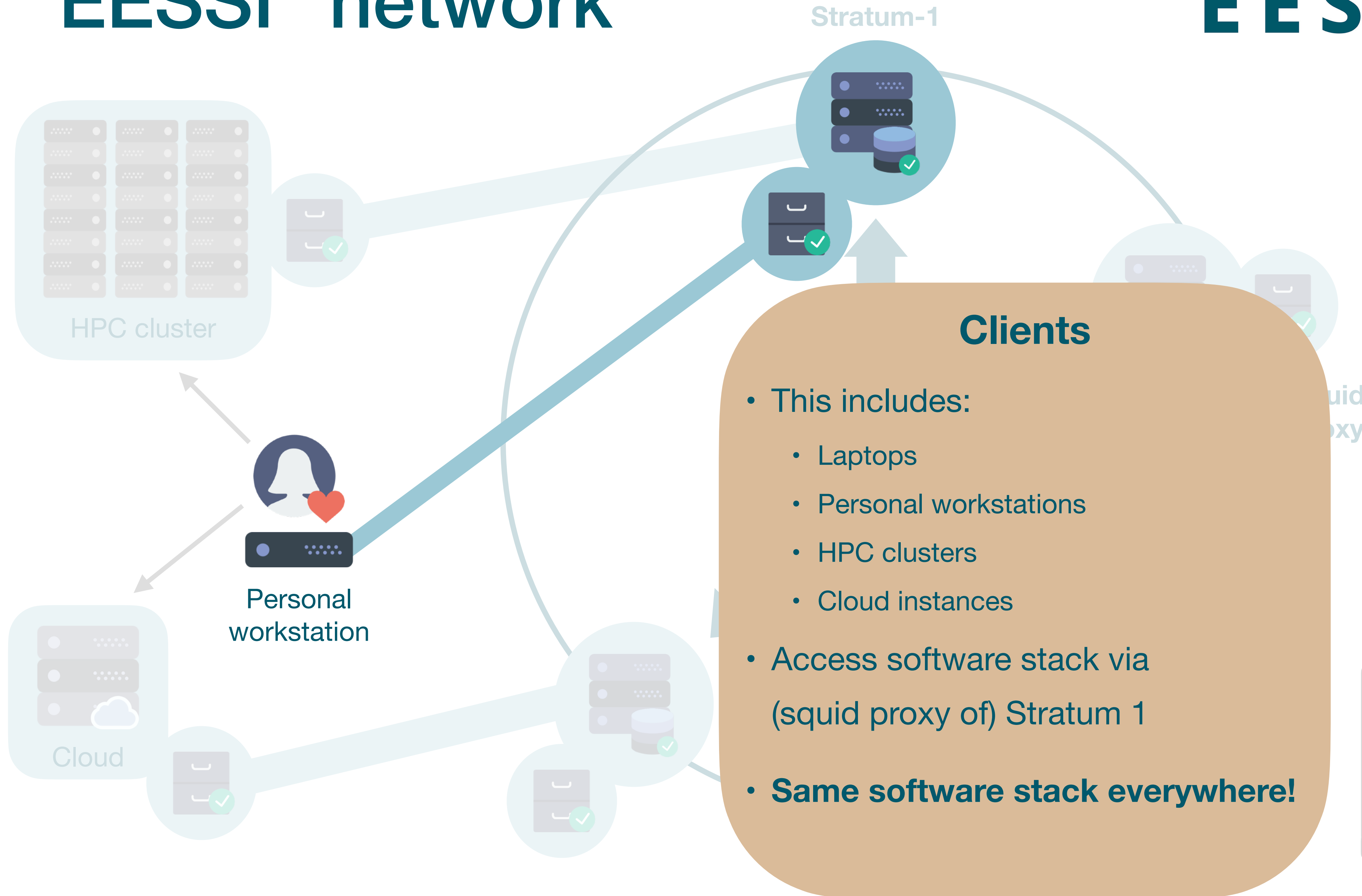
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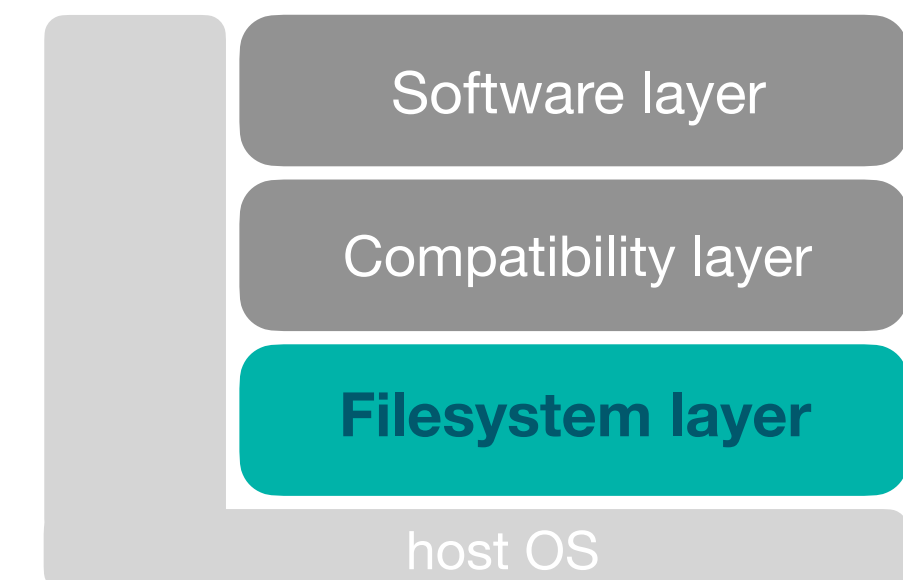
CernVM-FS



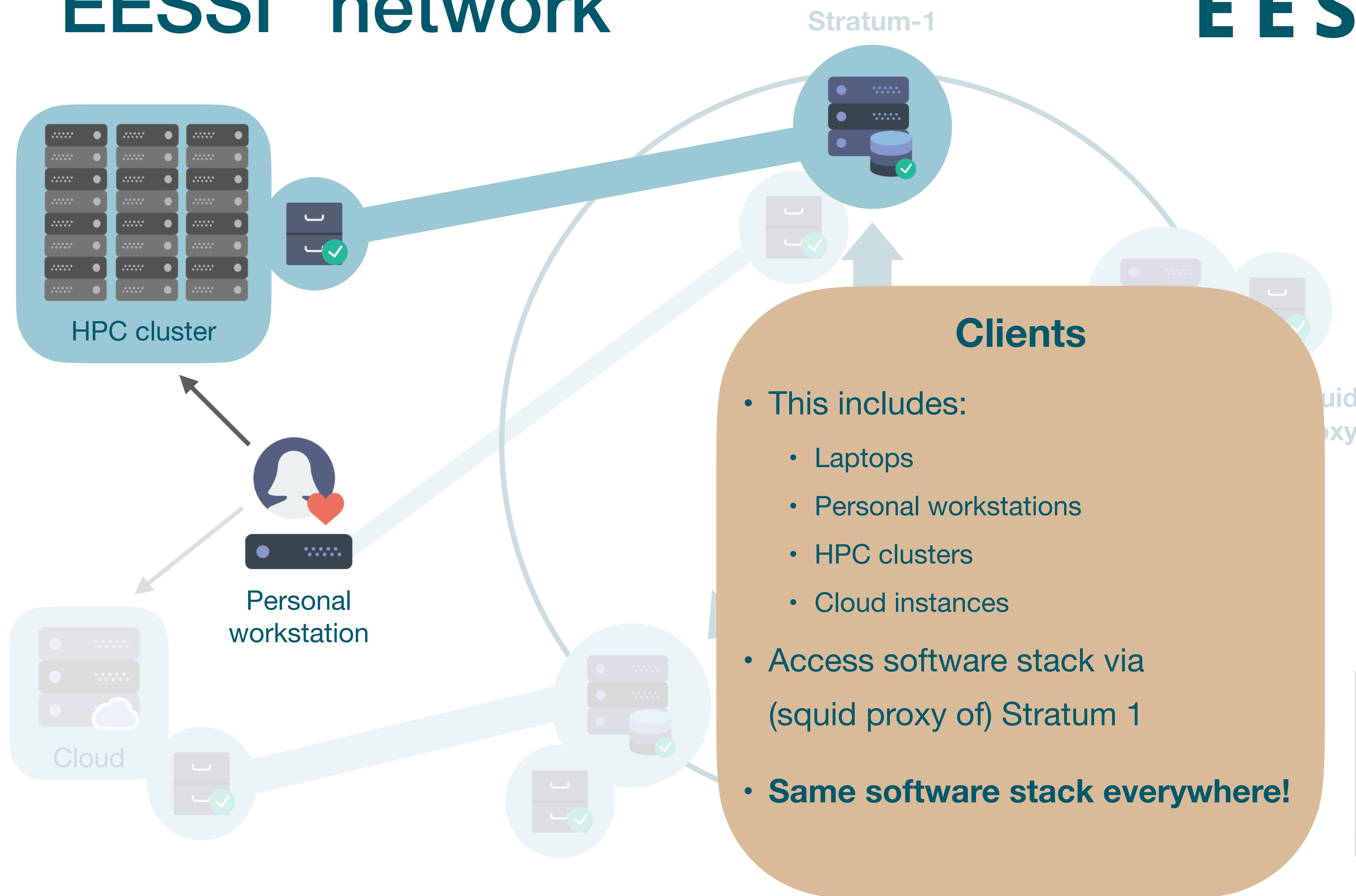
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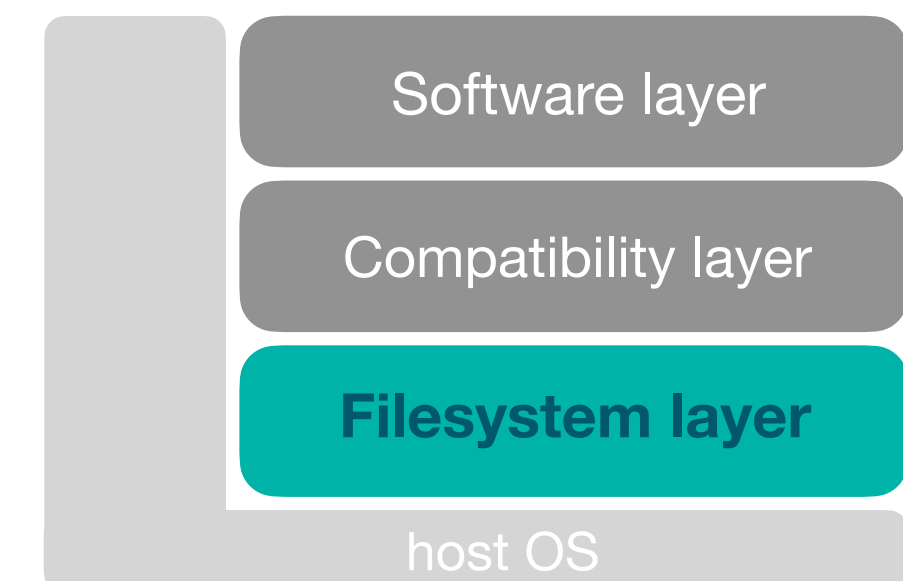


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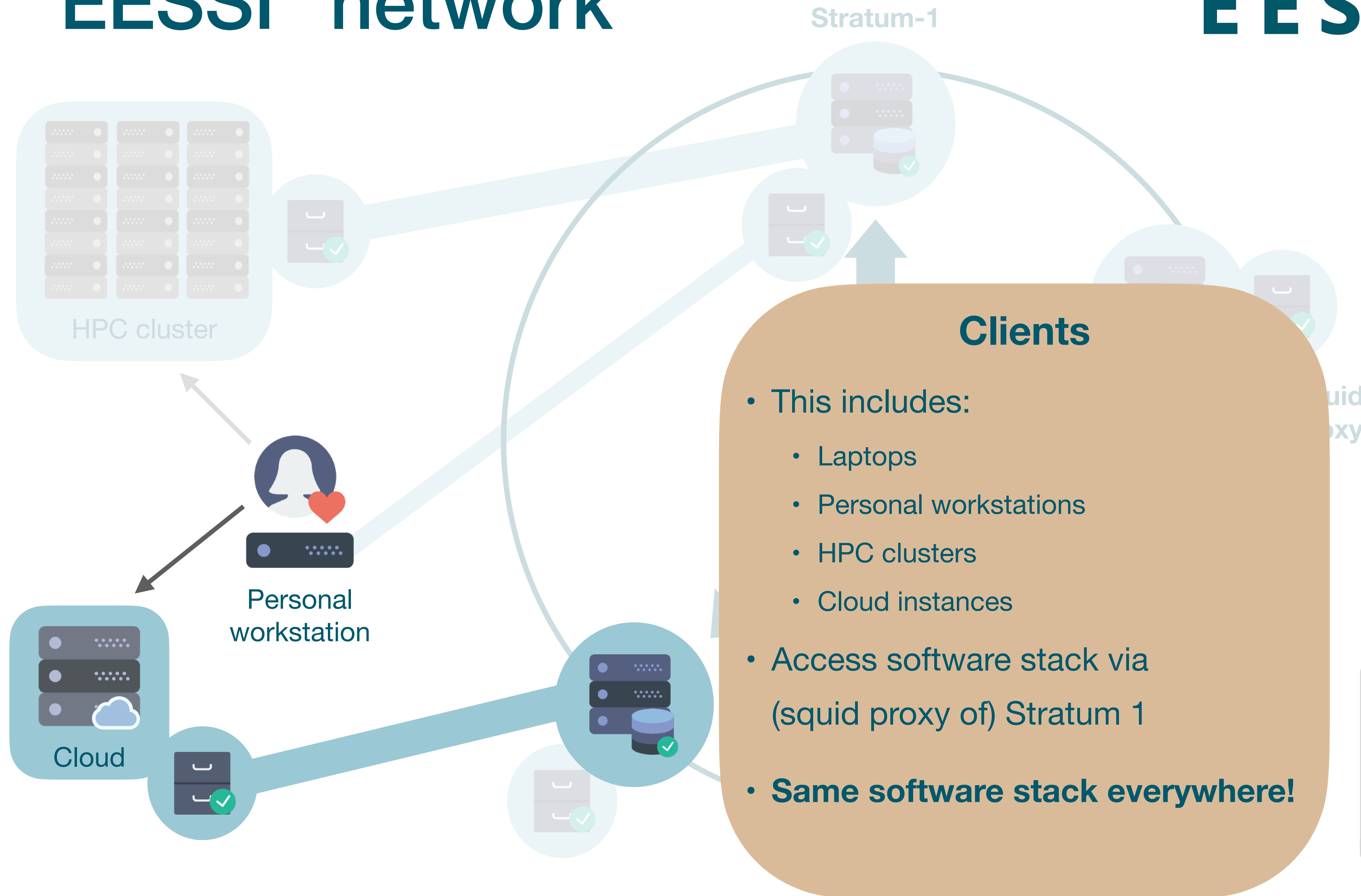


- This includes:
 - Laptops
 - Personal workstations
 - HPC clusters
 - Cloud instances
- Access software stack via (squid proxy of) Stratum 1
- **Same software stack everywhere!**

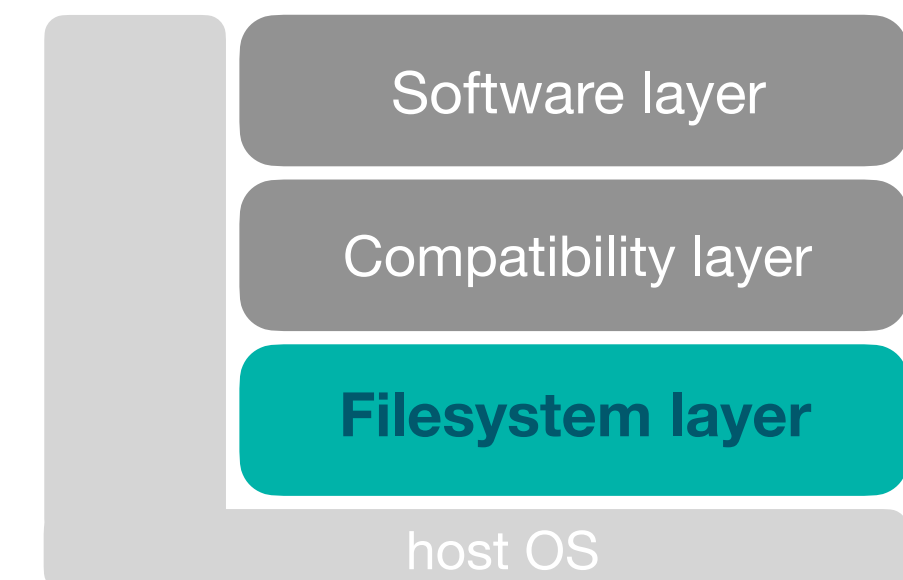
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