



# Running Web Applications on HPC using containers

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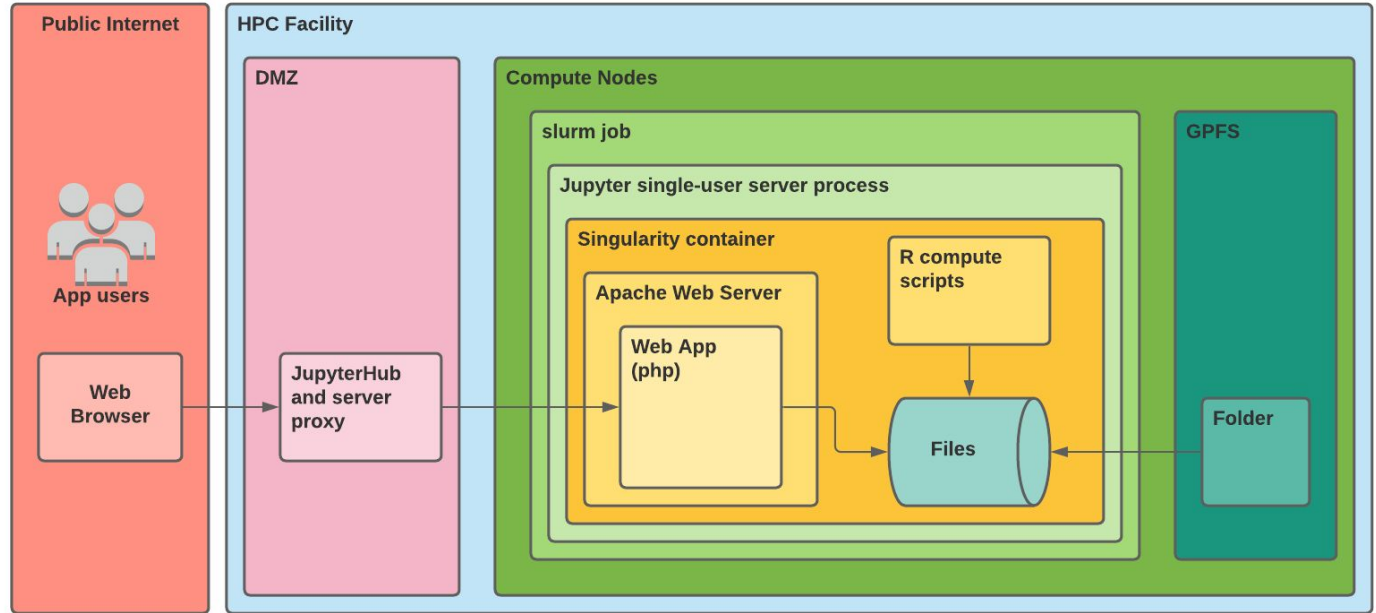
## What problem are we trying to solve?

- Hosting public facing web apps in a modern cloud environment is ubiquitous, not so common in an HPC environment
  - Easy to do in the cloud because:
    - Infrastructure designed to support this concept at scale from day 1
    - Huge community of practice, ecosystem of tools, DevOps
    - Standard service fully supported by commercial operators
  - Still hard to do for a Researcher on HPC because:
    - No standard HPC specific tooling/workflows/support to do this job so usually need to roll-your-own
    - HPC system architecture is rarely designed to specifically cater for this scenario
    - Security, availability, maintainability, complexity challenges
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## Who needs this?

- Q: Why bother at all to expose web apps on HPC when it is much easier and almost free to do so in the cloud?
- A: Web apps provide an easy way to submit jobs and access job results - this extends the audience for an experiment/analysis tool that requires a lot of CPU/Memory/GPU or has specific data sovereignty needs. Doing this in the commercial cloud would be too costly or geographically infeasible.

# What is our solution?



**Note:** Our aim was to use the set of available tools to deliver the solution in a timely manner, re-designing the infrastructure for this purpose was out of scope

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Concrete example  
and story - running  
a web app on the  
HPC using  
containers

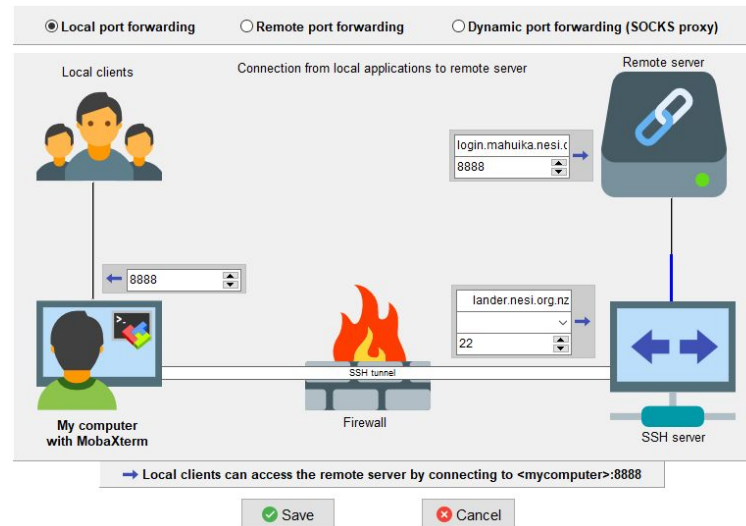
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- Working with Basharat Bhat (University of Otago and Genomics Aotearoa)
    - He is developing a web app for running an Epigenome Wide Association Study (EWAS) pipeline
    - Requires lots of memory to run (>100GB)
  - Docker and singularity images for portable web app:  
[https://github.com/GenomicsAotearoa/EWAS\\_DAP](https://github.com/GenomicsAotearoa/EWAS_DAP)
  - Looked into the options for running the web app on NeSI

This work was done through NeSI's Consultancy Service:  
<https://www.nesi.org.nz/services/consultancy>

Contact us if interested to find out how we can support your project.

## Concrete example and story

- Initial solution (SSH based)
  - Researcher SSH's into the cluster, edits Slurm script, chooses a port and submits job
  - Then sets up SSH tunnel on chosen port from their machine to cluster



## Concrete example and story

- Better solution (Jupyter Server Proxy based)
  - Connect to Jupyter - <https://jupyter.nesi.org.nz>
  - Request required resources (memory, wall time, ...)
  - From the JupyterLab launcher, click the “EWAS” button to launch the web app (runs within the resources you requested above)
  - Completely browser based, no SSH or terminal required



# Demo

- Use Jupyter Server Proxy to run the web app via NeSI's JupyterLab service
  - Simple to configure
  - Requires a script to run web app
  - Accept options to specify port and base\_url

```
jupyter_notebook_config.py
1 c.ServerProxy.servers = {
2   'EWASP': {
3     'command': ['/path/to/run_ewas.sh', '{port}', '{base_url}'],
4     'timeout': 10,
5   },
6 }
```

Demo - <https://jupyter.nesi.org.nz>



- Containers
  - Docker image bundling the php web app, web server, compute R scripts and dependent libraries
  - Singularity image built from the Docker image
  - Data mapped from HPC file system via container volumes
  - Various hacks in the application to enable serving through tunnel/proxy (i.e. relative paths)
- Jupyter Proxy - <https://github.com/jupyterhub/jupyter-server-proxy>
  - Allows the user to proxy to a server running somewhere on the cluster (can be within the same node/allocation or elsewhere)
  - Can also start/manage the process if required (shows up as a button in the launcher next to the kernels)
  - Workaround for not being able to expose services running on the HPC to the outside world

## What were the challenges?

- Can't use docker on the HPC
  - Still wanted a docker version for portability
  - Convert docker to singularity (workarounds for non-root)
- Can't easily expose to the outside world
  - SSH is the usual way to access the HPC
  - Requires multi-hop SSH tunnelling to expose web app
- HPC has a scheduler / queueing system
  - Not ideal for interactive use, may have to wait for resources
- Complex to deploy/use
  - Can't build on the HPC - need to build locally or in the cloud
  - Many R packages need to be built from source - time consuming build process
  - Web app relative paths don't work out of the box when running through a proxy/tunnel

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## Q&A

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Questions?

[support@nesi.org.nz](mailto:support@nesi.org.nz)

<https://www.nesi.org.nz/apply>



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