

# The Genomics Virtual Laboratory 4.0

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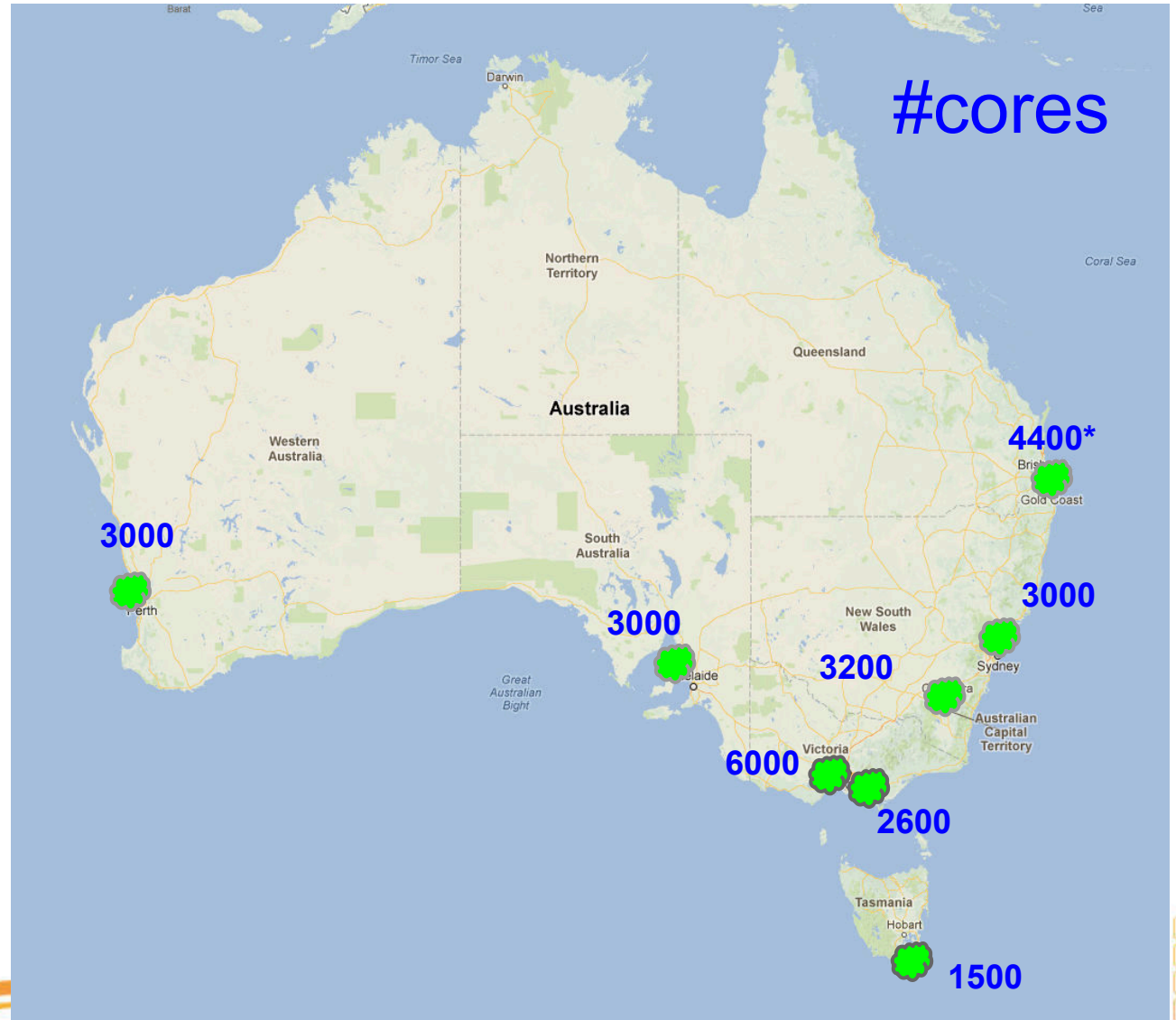


**Government funded, free for researchers**

**30,000 cores + co-located object and NFS stores**

**OpenStack based**

**~\$100m USD**



1. Goto [www.genome.edu.au](http://www.genome.edu.au)
2. Use the public galaxy servers
3. or get your own private instance
  - a. Click Get -> Launch Your Own GVL
  - b. Launch!



# GVL: Design principles

<b>Criteria</b>	<b>Design Implication</b>
<b><i>Accessible</i></b>	Minimal client-side requirements
<b><i>Reproducible</i></b>	Workflow support + software & tool management process
<b><i>Performance</i></b>	User-managed scaling of compute resources + high availability resources
<b><i>Flexible</i></b>	User configurable + administrable Multiple interaction modes
<b><i>Consistent</i></b>	Single platform from training to analysis
<b><i>Functional</i></b>	Pre-populated with suite of tools for common use cases + required reference data + visualisation options

# Resourcing

**Managed service: objective**



# Resourcing

Managed service: objective



A short time later...



# GVL: Philosophical assertion



# GVL: Philosophical assertion 2015





**GET**



**USE**



	<b>Personal GVL</b>	<b>Server GVL</b>	<b>Cluster GVL</b>
<i>Suitable for</i>	<b>Single user</b>	Single user Small group/lab	Large groups Institutions
<i>Storage</i>	<b>60GB</b>	100-5000GB	TBs
<i>Compute</i>	<b>2 cores</b>	8-64* cores	>50 cores
<i>Requires</i>	<b>NeCTAR account</b>	NeCTAR allocation: Compute and Volume storage	Large NeCTAR allocation of compute + user-provided fast storage
<i>Runs on</i>	<b>Any Research Cloud node</b>	RC nodes with volumes	RC nodes co-located with fast file system
<i>Setup</i>	<b><u>Automatic via website</u></b>	<b><u>Automatic via website</u></b>	Collaboration with GVL team
<i>Configuration</i>	<b>No configuration required</b>	Some configuration to tune analyses	Dedicated management

GVL: <http://genome.edu.au/>



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1. Flavours are pre-built variations of the GVL
2. Built by extending the base GVL ansible scripts (The GVL ansible scripts will internally include the Galaxy and Cloudman scripts)
3. Only modifications need to be implemented - minimising redundancy

Example: Microbial GVL

4. Made reproducible by defining exact component versions in launcher.

1. Each version of GVL has a fixed configuration so the software configuration is fully reproducible.
2. Since GVL 3.x, improvements have been made so that every single component is locked down.
3. Both automated testing and manual testing employed to make sure base functionality is correct.

