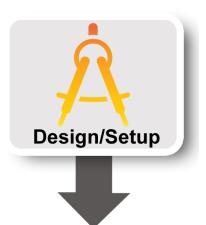


Getting Started with GENI -Part 2



ICDCS 2013 8 July 2013









Log into the GENI

Portal

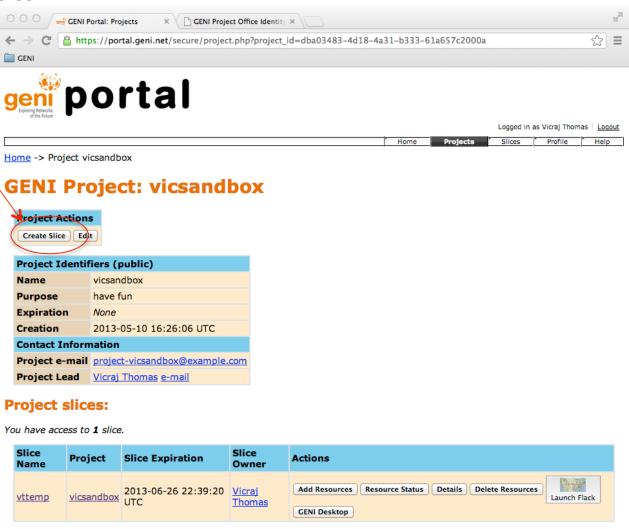




Log into the GENI

Portal

Create a slice



Expired Slices



https://portal.geni.net/secure/slice.php?slice_id=a163bd89-1522-46a5-b971-075489cb91fe

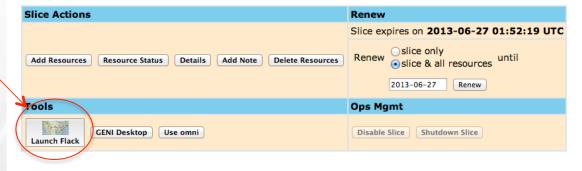
Geni Desktop! × V 🚟 GENI Portal: SII × V 🚟 GENI Key Mana × V 🚟 GENI Key Mana × V 📑 GENI Project O ×

Projects

- Log into the GEN GEN Portal
- Create a slice
- Launch the Flack experimenter tool GENI Slice: vttemp2



Home -> Project vicsandbox -> Slice vttemp2 Created slice vttemp2



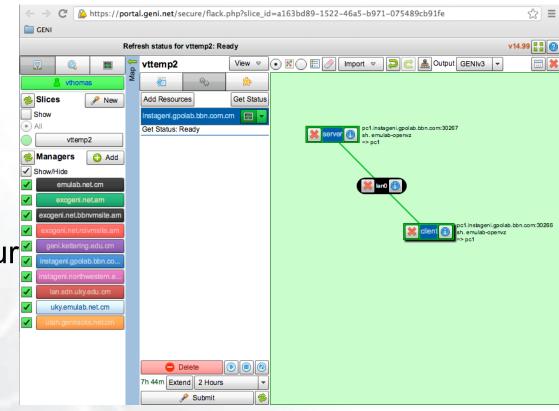
Confused? Look at the Portal Help or GENI Glossary.

Slice Status

Status Get All	Aggregate	Renew	Actions
not retrieved Get Status	GPO ExoGENI	Expires on not retrieved 2013-06-27 Renew	Resource Status Details Delete Resources
not retrieved	RENCI ExoGENI	Expires on not retrieved	Resource Status Details Delete Resources

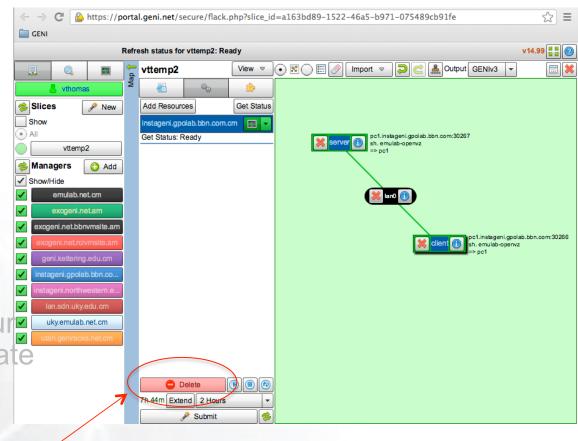


- Log into the GENI Portal
- Create a slice
- Launch the Flack experimenter tool
- Set up a simple experiment
 - Add resources to your slice from an aggregate
 - Use the resources in your slice



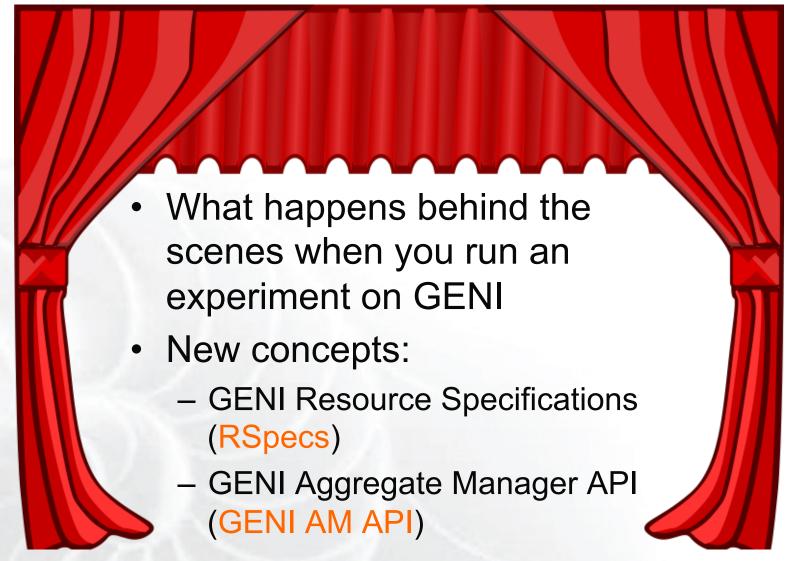


- Log into the GENI Portal
- Create a slice
- Launch the Flack experimenter tool
- Set up a simple experiment
 - Add resources to your slice from an aggregate
 - Use the resources in your slice
- Delete resources in your slice





In Part 2 You will Learn...



NIGENI Resource Specifications (RSpecs)

- RSpecs are XML documents that describe resources
 - VMs, links, etc.

RSpec for a virtual machine with one interface:

```
<?xml version="1.0" encoding="UTF-8"?>
<rspec type="request" xsi:schemaLocation="http://www.qeni.net/</pre>
resources/rspec/3 ... xmlns="http://www.geni.net/resources/rspec/3">
    <node client id="server" component manager id="urn:publicid:IDN</pre>
+instageni.gpolab.bbn.com+authority+cm">
       <sliver type name="emulab-openvz"/>
       <interface client id="server:if0"> </interface>
    </node>
</rspec>
```



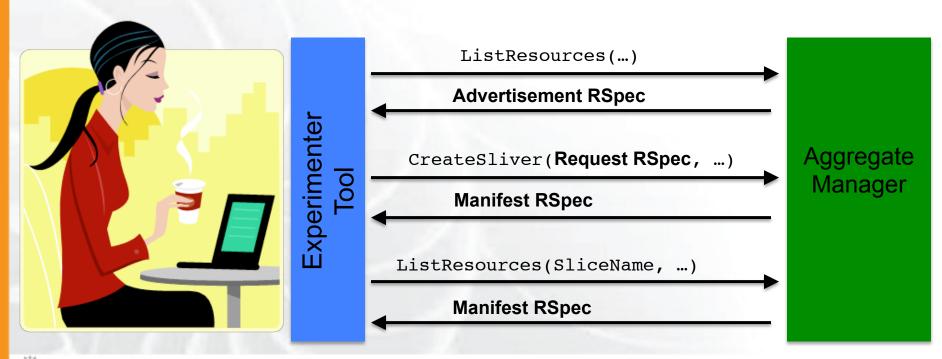


- RSpec documents are exchanged by experimenter tools (e.g. Flack) and aggregates
 - Aggregates use RSpecs to describe what they have –
 Advertisement RSpecs
 - Experimenters use RSpecs to describe the resources they want – Request RSpecs
 - Aggregates use RSpecs to describe the resources allocated to an experimenter – Manifest RSpecs



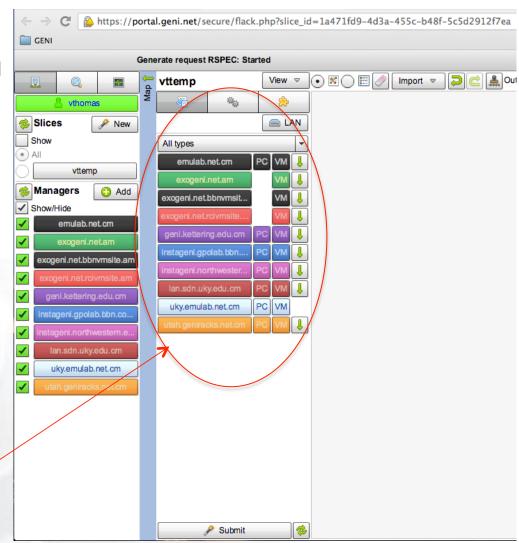
The GENI AM API

 Experimenter tools and aggregates talk to each other using the GENI Aggregate Manager API (GENI AM API)



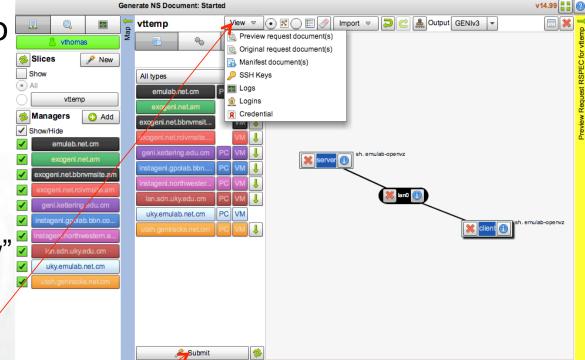


- Flack calls ListResources on all aggregates it knows about
- Aggregates send back advertisement **RSpecs**
- Flack uses information in the advertisements to populate its palette of resources





- When you drag and drop resources on to the Flack canvas, it creates request **RSpecs** for these resources
 - To view the request Rspec click on "View" and select "Preview request documents"
- When you click "Submit", Flack makes createSliver calls on the aggregates





Flack periodically calls sliverStatus on

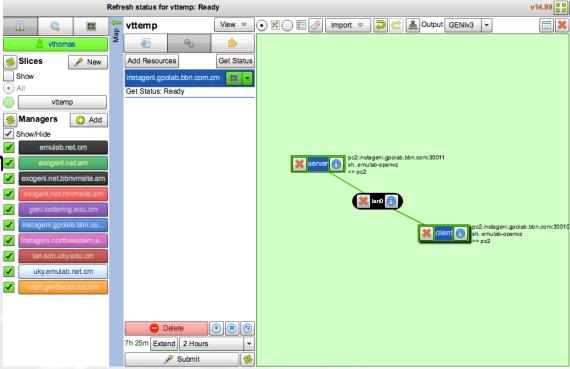
the aggregates to check on the status of

your request

When sliverStatus shows:

> Resources have been allocated, Flack turns its canvas yellow

 Resources are ready to use, Flack turns its canvas green



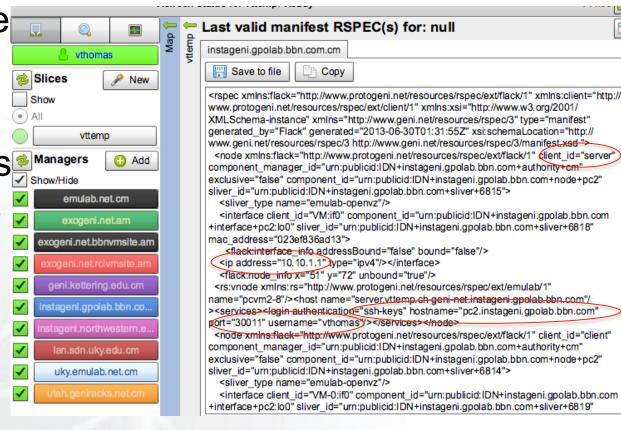
https://portal.geni.net/secure/flack.php?slice_id=1a471fd9-4d3a-455c-b48f-5c5d2912f7ea



listResources with a slice name returns a

manifest RSpec

- Manifest includes Managers names and ports used to ssh into **VMs**
 - Flack uses this information to help you log into your resources

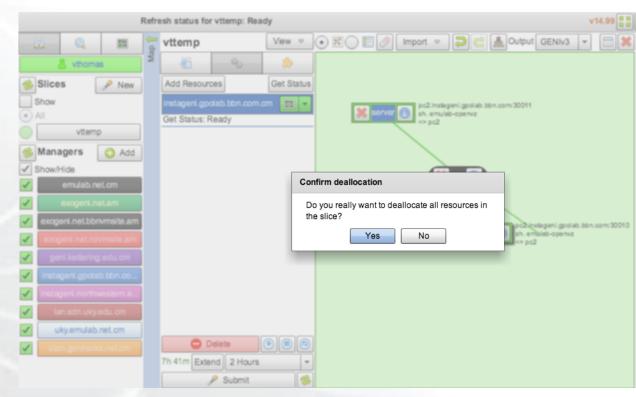




When you deleted your resources, Flack called

deleteSliver on

the aggregates





Hands-On Exercise

- Reinforce understanding of the new concepts by:
 - Viewing and editing RSpec documents
 - Making the AM API calls ourselves using the Omni experimenter tool



The Exercise

- Repeat the experiment from Part 1 by loading an RSpec into Flack
 - Instead of drawing the topology ourselves (saves time)
- Edit the RSpec using Flack but don't "submit"
- Save the request RSpec generated by Flack into a file
- Use Omni to make GENI AM API calls to send the request RSpec, check status of resources, etc.



GENI AM API Calls Used

- listresources: Get an advertisement rspec listing the resources at an aggregate
- createsliver: Request resources from an aggregate
- sliverstatus: Get status of resources allocated to a slice at an aggregate
- deletesliver: Delete resources allocated to a slice by an aggregate

For a complete list of GENI AM API calls see: http://groups.geni.net/geni/wiki/GAPI AM API



Omni: Resource Reservation tool

- A command line experimenter tool
- Useful for making AM API calls on aggregates
- Written in and scriptable from **Python**
- Works with aggregates that implement the GENI AM API
 - ProtoGENI, PlanetLab, OpenFlow, InstaGENI, **ExoGENI**

```
$ omni.py createsliver aliceslice myRSpec.xml
INFO: omni: Loading config file omni config
INFO:omni:Using control framework pgeni
INFO:omni:Slice urn:publicid:IDN+pgeni.gpolab.
          expires within 1 day on 2011-07-07
INFO:omni:Creating sliver(s) from rspec file
INFO:omni:Writing result of createsliver for
INFO:omni:Writing to 'aliceslice-manifest-rspe
INFO:omni:
INFO:omni: Completed createsliver:
  Options as run:
                aggregate: https://www.emulab.
                framework: pgeni
                native: True
  Args: createsliver aliceslice myRSpec.xml
  Result Summary: Slice urn:publicid:IDN+pgeni
Reserved resources on https://www.emulab.net/p
  Saved createsliver results to aliceslice-man
```

http://trac.gpolab.bbn.com/gcf/wiki/Omni





Omni Commands You Will Use

- omni.py —a aggregatename listresources
- omni.py —a aggregatename createsliver slicename requestRSpec
- omni.py —a aggregatename sliverstatus slicename
- omni.py —a aggregatename listresources slicename
- omni.py —a aggregatename deletesliver slicename

- A useful utility (distributed with Omni): readyToLogin.py
 - Gives you the ssh commands you need to log into your nodes
 - readyToLogin.py parses the output of sliverStatus to determine the hostname, portname and username for the ssh commands



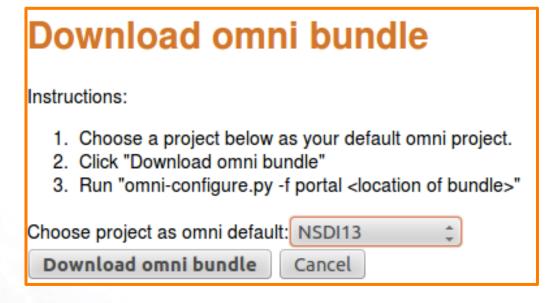
Configuring Omni

- Omni reads a configuration file omni config to:
 - Get usernames for accounts to be created on compute resources
 - Find locations of ssl certs and ssh key files
 - ssl certs are used to secure communication between Omni and the aggregates
 - ssh key pairs are used log into compute resources
 - Find standard nicknames for aggregates
 - E.g. you can refer to the InstaGENI rack at BBN as ig-bbn instead of https://boss.instageni.gpolab.bbn.com:12369/ protogeni/xmlrpc/am/2.0



Creating an Omni Config File

- Download the GENI bundle from the GENI **Portal**
- 2. Run the script omniconfigure.py
 - Distributed with Omni
 - Already installed on your virtual machine



geni@NSDI13-Tutorials:~\$> omni-configure.py -f portal



Watch Instructor do the Exercise





Wrapping Up...

- By the end of this tutorial you should:
 - Feel comfortable running simple experiments on GENI
 - Have a basic understanding of how GENI works
- Later tutorials may skip some of these basic steps to focus on new material
 - You may be given an RSpec to use rather than have you create one
 - You may use slices that have already been created and resources added to them

For a description of the GENI concepts you have learned, see: http://groups.geni.net/geni/wiki/GENIConcepts



Try it on Your Own...

- You should have a printout with detailed instructions
- Online instructions:

http://groups.geni.net/geni/wiki/ GENIExperimenter/Tutorials/PortalOmniExample