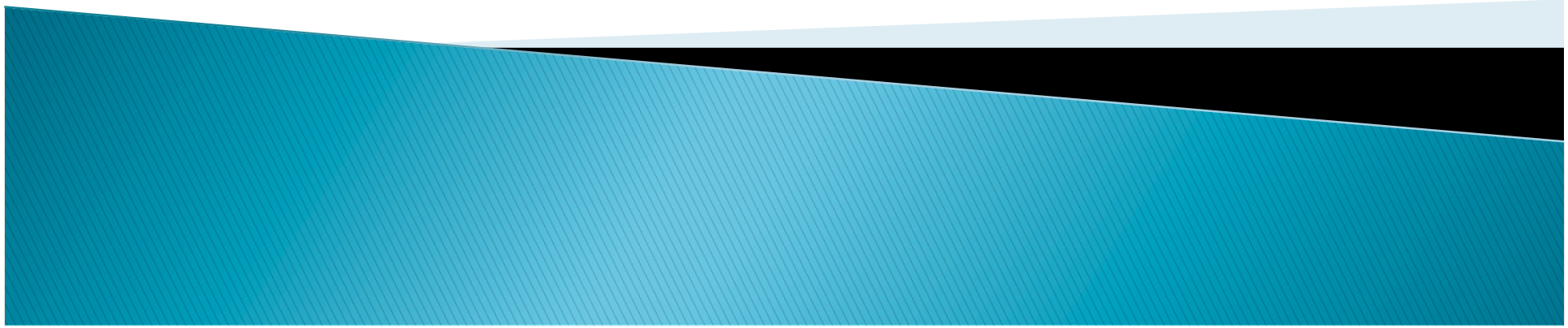
ExoGENI/GIMI Tutorial
Part 1: ExoGENI

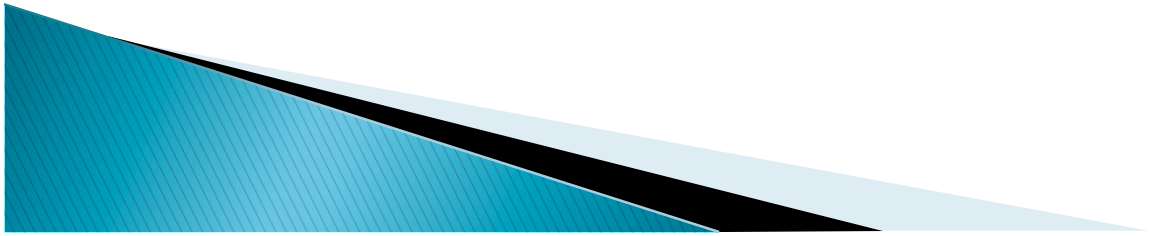
Ilia Baldine ibaldin@renci.org

Anirban Mandal, Yufeng Xin



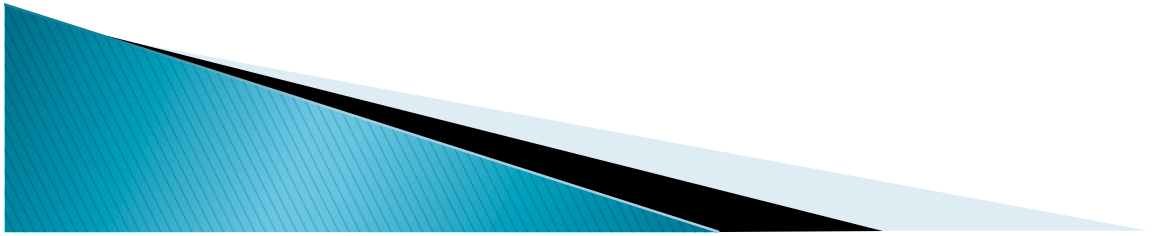
Tutorial sections

- Configure environment
- ExoGENI and Orca Overview
- Creating slices with Omni and GENI AM API
- Flukes Overview
- Creating slices with Flukes
- Tutorial page:
  - http://groups.geni.net/geni/wiki/ORCAExoGENITutorial
  - Please open in your browser
  - Please open the presentation



Configuring Environment

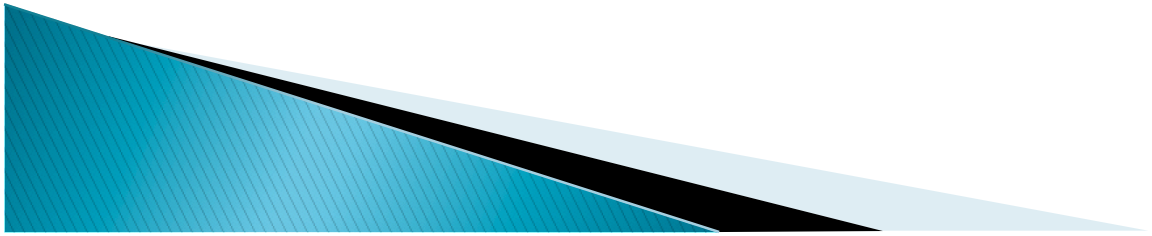
- All user properties are under $HOME/.flukes.properties – it is a text file
  - Edit $HOME/.flukes.properties
    - Open in an editor and replace *EVERY* occurrence of XX with your index (including leading zero)
    - Update the orca.xmlrpc.url property
  - Inspecting keystore file (make note of key alias ('tutorialXX')
    - $ cd Tutorials/GIMI/gimiXX
    - $ keytool -list -keystore ssh/gimi01.jks
    - NOTE: your key name is gimiXX, and your key and keystore password is g5C7r#XX



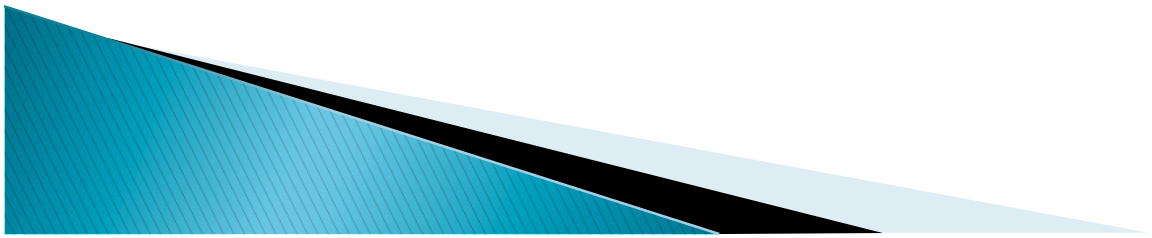
Rack binding

- Today's tutorial will use 3 separate AMs

- Users 01-05 -> RCI
  - https://rci-hn.exogeni.net:11443/orca/xmlrpc
- Users 06-10 -> BBN
  - https://bbn-hn.exogeni.net:11443/orca/xmlrpc
- Users 11-20 -> NICTA
  - https://nicta-hn.exogeni.net:11443/orca/xmlrpc



Section: ExoGENI and ORCA Overview



ExoGENI Testbed

- 14 GPO-funded racks
  - Partnership between RENCI, Duke and IBM
  - IBM x3650 M3/M4 servers
    - 48G RAM
    - Dual-socket 8-core Intel X5650 2.66Ghz CPU
    - 10G dual-port Chelseo adapter
  - BNT 8264 10G/40G OpenFlow switch
  - DS3512 6TB sliverable storage
- Each rack is a small networked cloud
  - OpenStack- and xCAT based
  - EC2 nomenclature for VM node sizes (m1.small, m1.large etc)
  - Baremetal node provisioning
  - Interconnected by combination of dynamic and static L2 circuits through regionals and national backbones
- http://wiki.exogeni.net

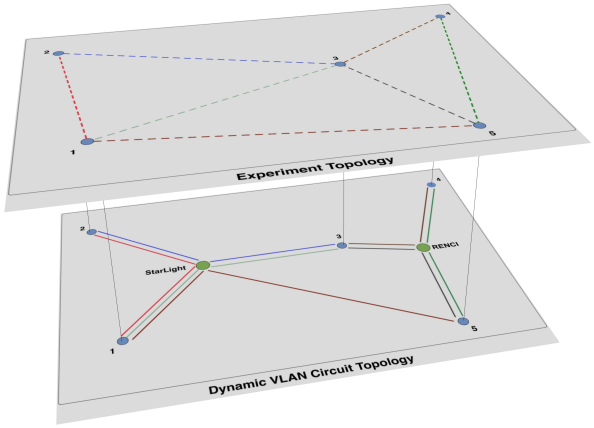
ExoGENI Status

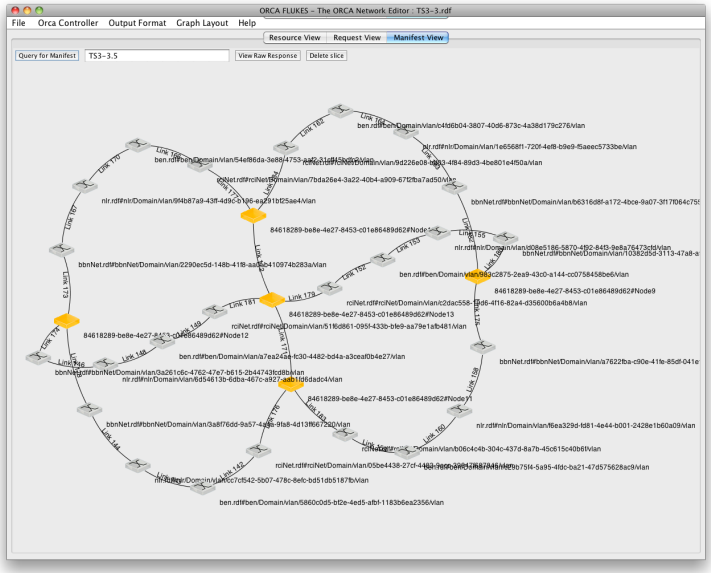
- 2 new racks deployed
  - RENCI and GPO
- 2 older existing racks
  - Duke and UNC
- 2 more racks coming
  - FIU and UH
- Partner racks
  - NICTA
  - U of Alaska Fairbanks

- Connected via BEN (http://ben.renci.org), LEARN and NLR FrameNet, (eventually I2)

ExoGENI slice isolation

- Strong isolation is the goal
- Compute instances are KVM based and get a dedicated number of cores (ExoGENI does not over-provision cores)
  - Caveat: currently all instances get 1 core (different RAM and disk).
- VLANs are the basis of connectivity
  - VLANs can be best effort or bandwidth-provisioned (within and between racks)
  - Caveat: current hardware in the racks allows best-effort VLANs only – will be remedied by Fall 2012 with support from the vendor





ORCA Overview

- Originally developed by Jeff Chase and his students at Duke
- Funded as Control Framework Candidate for GENI
  - Jointly developed by RENCI and Duke for GENI since 2008.
- A federation of networked clouds with a variety of interfaces
  - Native ORCA
  - GENI AM API
- Unique feature of ExoGENI: experimenter can
  - Operate on individual racks as independent aggregates
  - Operate on entire testbed and link racks together using ExoSM

ORCA deployment in ExoGENI

- Each rack runs its own Orca actor called the 'SM' that presents as GENI AM and exposes
  - ORCA native API
  - GENI AM API
- Rack-local SM can only create slices with **resources within that rack (virtual machines, baremetal nodes and vlans)**
- 'ExoSM' has global visibility
  - Has access to a fraction of resources **resources in all racks**
  - Has access to **network backbone resources for stitching topologies between racks**
- ExoSM
  - https://geni.renci.org:11443/orca/xmlrpc
- Rack SMs:
  - RENCI Rack: https://rci-hn.exogeni.net:11443/orca/xmlrpc
  - BBN Rack: https://bbn-hn.exogeni.net:11443/orca/xmlrpc
  - NICTA Rack: https://nicta-hn.exogeni.net:11443/orca/xmlrpc

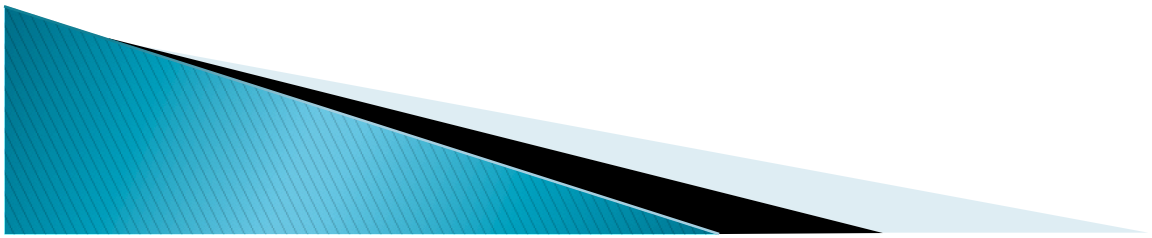
How are resources split?

- Resources in each rack are split between rack SM and ExoSM.
- Currently the split is 50/50 for VMs and internal VLANs between rack SM and ExoSM
- Baremetal nodes are usable only by ExoSM
- Stitching links from regional and national providers are delegated to ExoSM only.
- If your experiment is rack-local, you can always choose a specific rack SM to run your experiment
- If your experiment involves resources from multiple racks you must provision via ExoSM

How do I use it?

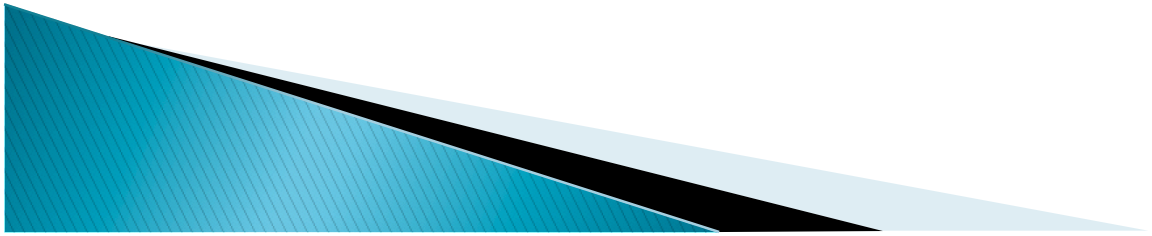
- Request credentials through the GPO
- Build your own VM Image [optional]
  - https://geni-orca.renci.org/trac/wiki/neuca-images
- Define topology and submit request
  - Use either Omni or Flukes
- Maximum slice lifetime is 2 weeks

Section: Creating slices with Omni



Creating slices with Omni and GENI AM API

- Use the RSpec file linked to this tutorial webpage:
  - http://groups.geni.net/geni/wiki/ORCAExoGENITutorial



Sample simple RSpec

```
<?xml version="1.0" encoding="UTF-8"?>
<rspec type="request"
      xsi:schemaLocation="http://www.geni.net/resources/rspec/3
                          http://www.geni.net/resources/rspec/3/request.xsd
                         http://www.protogeni.net/resources/rspec/ext/shared-vlan/1
                         http://www.protogeni.net/resources/rspec/ext/shared-vlan/1/request.xsd"
      xmlns:xsi="http://www.w3.org/2001/XMLSchema-instance"
      xmlns:s="http://www.protogeni.net/resources/rspec/ext/shared-vlan/1"
      xmlns=http://www.geni.net/resources/rspec/3>
<node client_id="geni1" component_manager_id="urn:publicid:IDN+bbnvmsite+authority+cm">
<sliver_type name="m1.small">
<disk_image name="http://geni-images.renci.org/images/standard/debian/debian-squeeze-amd64-neuca-2g.zfilesystem.sparse.v0.2.xml"
      version="397c431cb9249e1f361484b08674bc3381455bb9" />
</sliver_type>
<interface client_id="geni1:if0">
<ip address="172.16.2.1" netmask="255.255.255.0" />
</interface>
</node>
<node client_id="geni2" component_manager_id="urn:publicid:IDN+bbnvmsite+authority+cm">
<sliver_type name="m1.small">
<disk_image name="http://geni-images.renci.org/images/standard/debian/debian-squeeze-amd64-neuca-2g.zfilesystem.sparse.v0.2.xml"
      version="397c431cb9249e1f361484b08674bc3381455bb9" />
</sliver_type>
<interface client_id="geni2:if0" >
  <ip address="172.16.2.2" netmask="255.255.255.0" />
 </interface>
</node>
<link client_id="local">
  <interface_ref client_id="geni1:if0" />
  <interface_ref client_id="geni2:if0" />
</link>
</rspec>
```

Issuing OMNI commands (1/2)

- Look at 'getVersion()' output
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc getversion
- Download request RSpec from tutorial webpage
- Create a slice with GPO SA
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc createslice orcav2-test3
- CreateSliver with given RSpec
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc -n createsliver orcav2-test3 two-node.rspec

Issuing OMNI commands (2/2)

- Query sliver status
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc sliverstatus orcav2-test3
- List resources within the slice (IP addresses)
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc listresources orcav2-test3
- Login to nodes
  - ssh -i ssh/gimiXX_key root@<ip address>
- Delete sliver
  - omni.py -c omni_config -a https://geni.renci.org:11443/orca/xmlrpc deletesliver orcav2-test3

RSpec conventions

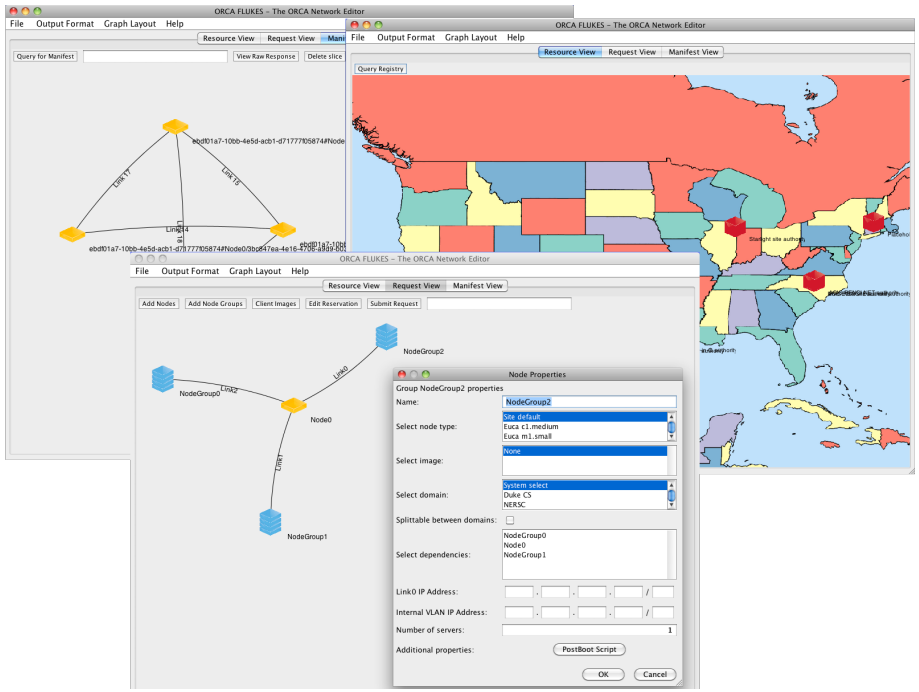
- *Install* and *execute* service tags are respected
  - tar.gz, tar.Z, tar.bz, deb, rpm and zip recognized
- Domain binding is possible by specifying component id or component manager id

```
<node component_id="urn:publicid:IDN+uncvmsite+node
  +vm"
      component_manager_id="urn:publicid:IDN+uncvmsite
  +authority+cm"
      client_id="pc175"
      exclusive="true">
   <sliver_type name="raw-pc" />
</node>
```

- See https://geni-orca.renci.org/trac/wiki/orca-and-rspec for updated information

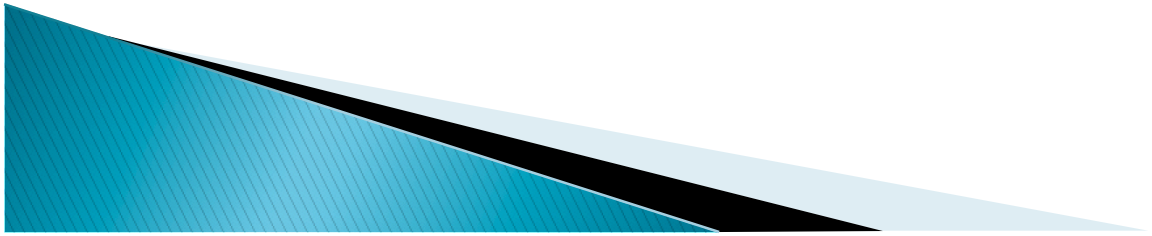
Section: Flukes Overview

- Graphical tool for creating and managing slice topologies in ORCA
  - JAVA (JNLP)



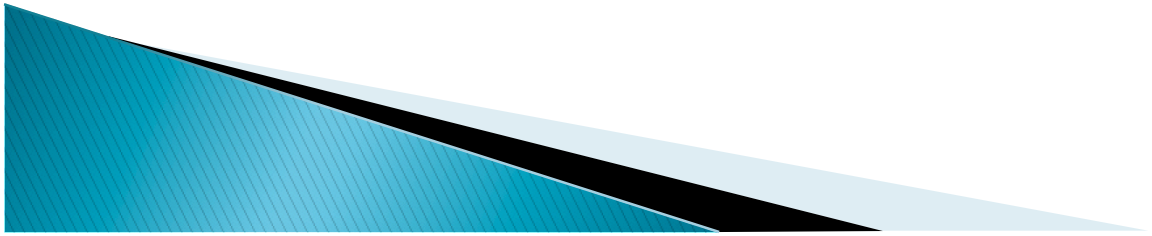
Section Overview

- Configuring Flukes prior to launch
- Launching Flukes
  - GUI Overview
  - Nodes, NodeGroups and Link parameters
  - Node-level vs. reservation level options
- Building slice request topologies
- Launching slice requests
- Inspecting slice manifests
- Logging into nodes in the slice
- NEuca-py tools



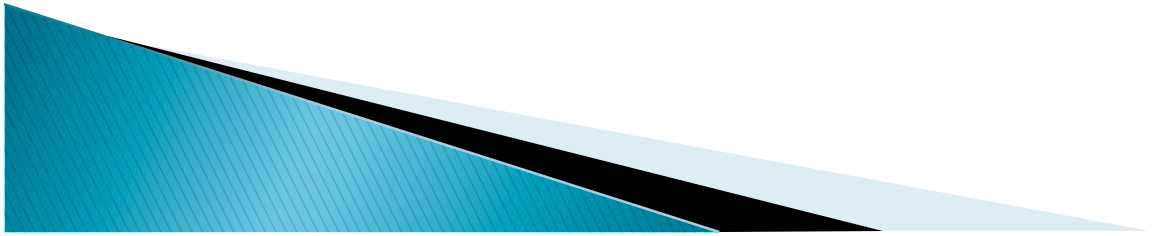
Launch Flukes!

- Double-click Flukes icon on your desktop
- Permanent stable version link
  - http://geni-images.renci.org/webstart/flukes.jnlp
- Can I use Flukes outside of ExoGENI?
  - No. Flukes uses semantic web mechanisms (RDF and OWL) to describe resources that is only compatible with ORCA and ExoGENI.



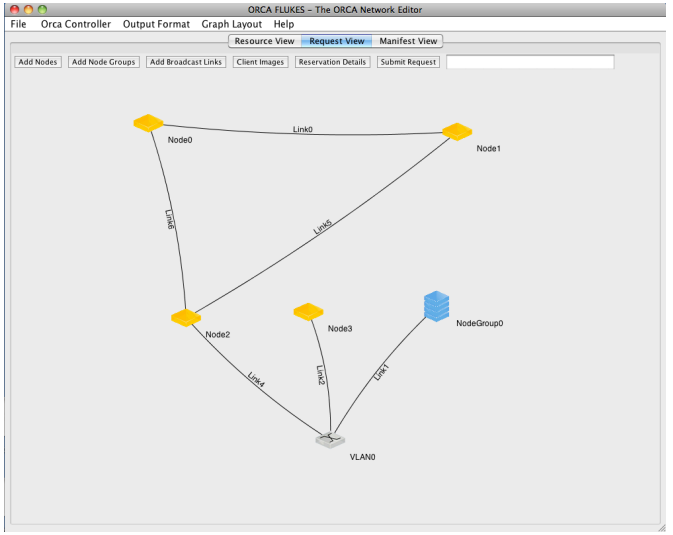
GUI Overview

- Tabs
  - Resources, Request, Manifest
- Menus
  - Current properties
  - Overwriting properties ($HOME/.flukes.properties)
- Mouse modes
- Buttons
- Adding nodes, nodegroups and links



Node and Link parameters

- Create a single node
- Right-click on the Node
  - Look at properties
  - Edit properties
    - Node type (size)
    - VM image
    - Domain (binding)
    - PostBoot script



- Create another node, link the two together
- Right-click and open properties again
  - Specify IP address on the link
  - Node functional dependencies
- Right-click on links
  - Inspect and edit link properties
  - Note only bandwidth is currently respected (and not everywhere due to hardware limitations)
- Broadcast links
- Specifying vlan tags
  - Only ‘special’ shared tags can be specified

NodeGroup parameters

- Create a single unattached node group
- Right-click to inspect and edit properties
  - Group sizes
  - Splittable groups

Nodes and NodeGroups

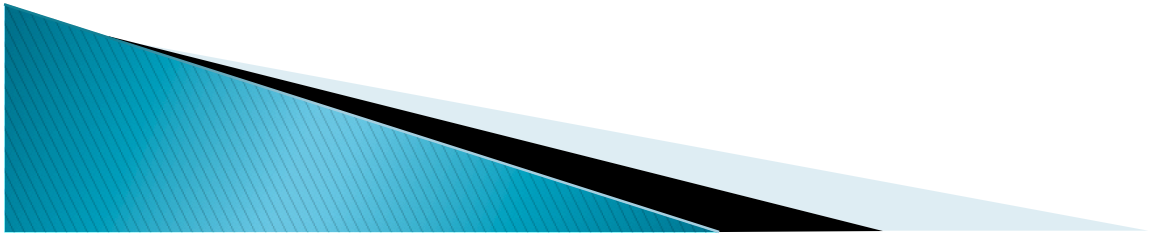
- A Node is an individual compute element
  - Typically a VM or a hardware node
  - IP address(es) on links, size, image, site binding, post boot script
  - Can I control management IP address assignment? NO!
- A NodeGroup is a group of identically configured nodes
  - A lot like a node except
    - PostBoot script is templated using Velocity template engine
      - https://geni-orca.renci.org/trac/wiki/flukes
    - IP address assignment is semi-automatic (starting with a user-specified address)
    - Node groups can be splittable between sites

What do I get when I …?

- Create a standalone node?
  - You get a single compute element at one of the sites with a single network interface to the management network through which you can SSH into the node
  - Management interface is always eth0
- Connect two nodes together?
  - You get two compute elements each with two network interfaces – one for management access and one for the link between two nodes.
  - User-controlled interfaces start with eth1
  - You can control IP address assignment on the interfaces linking the two nodes (suggested range: 172.16.0.0/16)

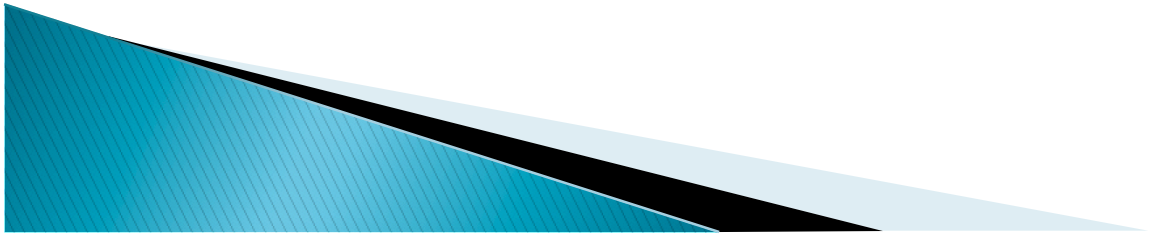
What do I get when I …?

- Create a standalone NodeGroup?
  - You get some number of nodes (specified in the group size) each with a single interface to the management network (eth0)
  - Nodes typically will be within the same rack
  - If node group is marked splittable nodes may be split across sites
- Connect a node group to a node or another node group?
  - All nodes within the group and the adjacent node (or all nodes in both groups) have interfaces on a common VLAN. They also have management interfaces (eth0)
  - IP address is specified similarly to private VLAN
  - Beware of address clashes! (i.e. here is a piece of rope, feel free to shoot yourself in the foot)



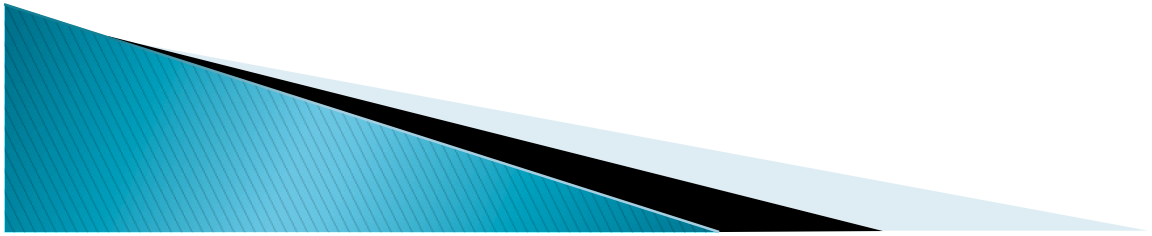
Can I tell which interface in the node will be eth1, eth2 etc?

- No, nor should you need to. Interfaces are identified by links they belong to.
- Note that different OSs name interfaces differently



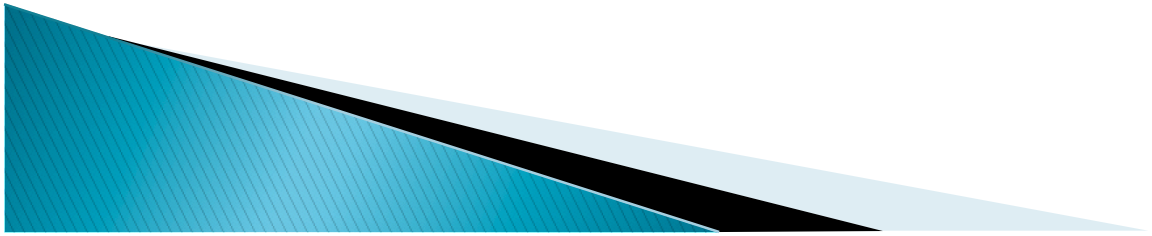
Node-level vs. Reservation-level options

- Reservation-level options overwrite node-level options
  - VM image
  - Domain binding
- OpenFlow slice parameters can currently only be specified at reservation level

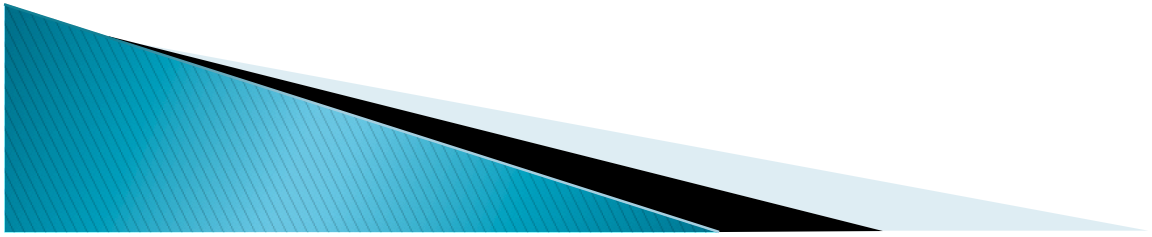


Domain binding

- Unbound requests are automatically bound to domains with available resources.
  - This depends on the visibility of the SM.
  - ExoSM can bind to any rack
  - Rack SM will always bind to its rack
- Bound requests are honored if resources are available
  - To create an inter-rack request, bind some of the nodes in it to one rack, and others to another
  - Can only be done via ExoSM!



Section: Creating slices with Flukes



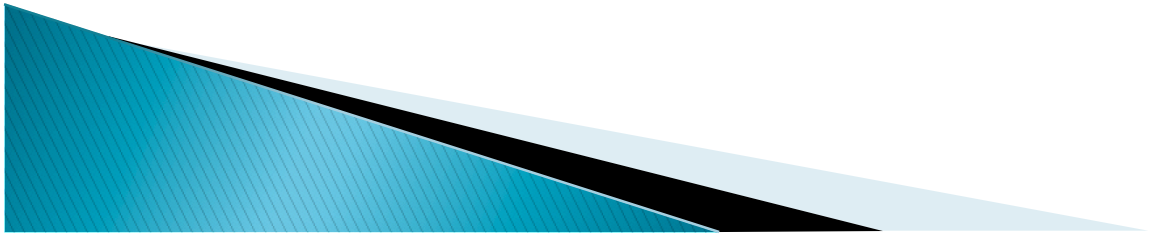
Launching a slice

- Retrieve the RDF request file:
  - cd ~/Tutorials/GIMI/gimiXX
  - wget http://emmy9.casa.umass.edu/RDFs/gimiXX.rdf

- Click 'File | Load Request' to load the request
- Inspect the topology, node properties and post boot scripts

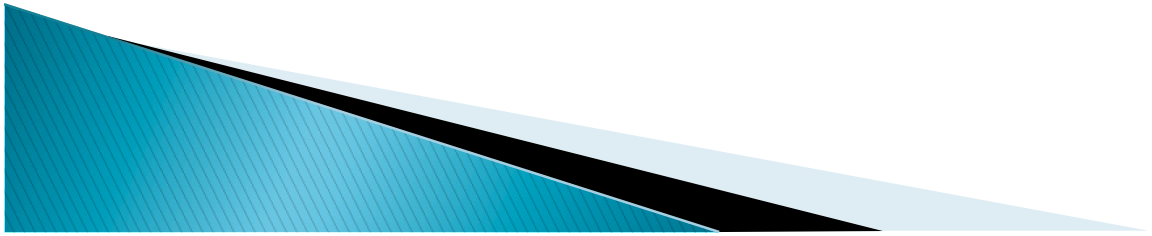
Launch a slice

- These will be unbound slices
  - We will let ORCA select sites with available resources
- Specify slice duration (click ‘Edit Reservation’)
- Fill in slice name (must be unique)
- Click ‘Submit Request’
  - Type in the alias of the key in the keystore (‘tutorialXX’)
  - Type in the password (‘tut0rialXX’)
  - ‘OK’
- Inspect the output window
  - Mainly a debugging tool. Will go away in the future.



Inspect slice manifest

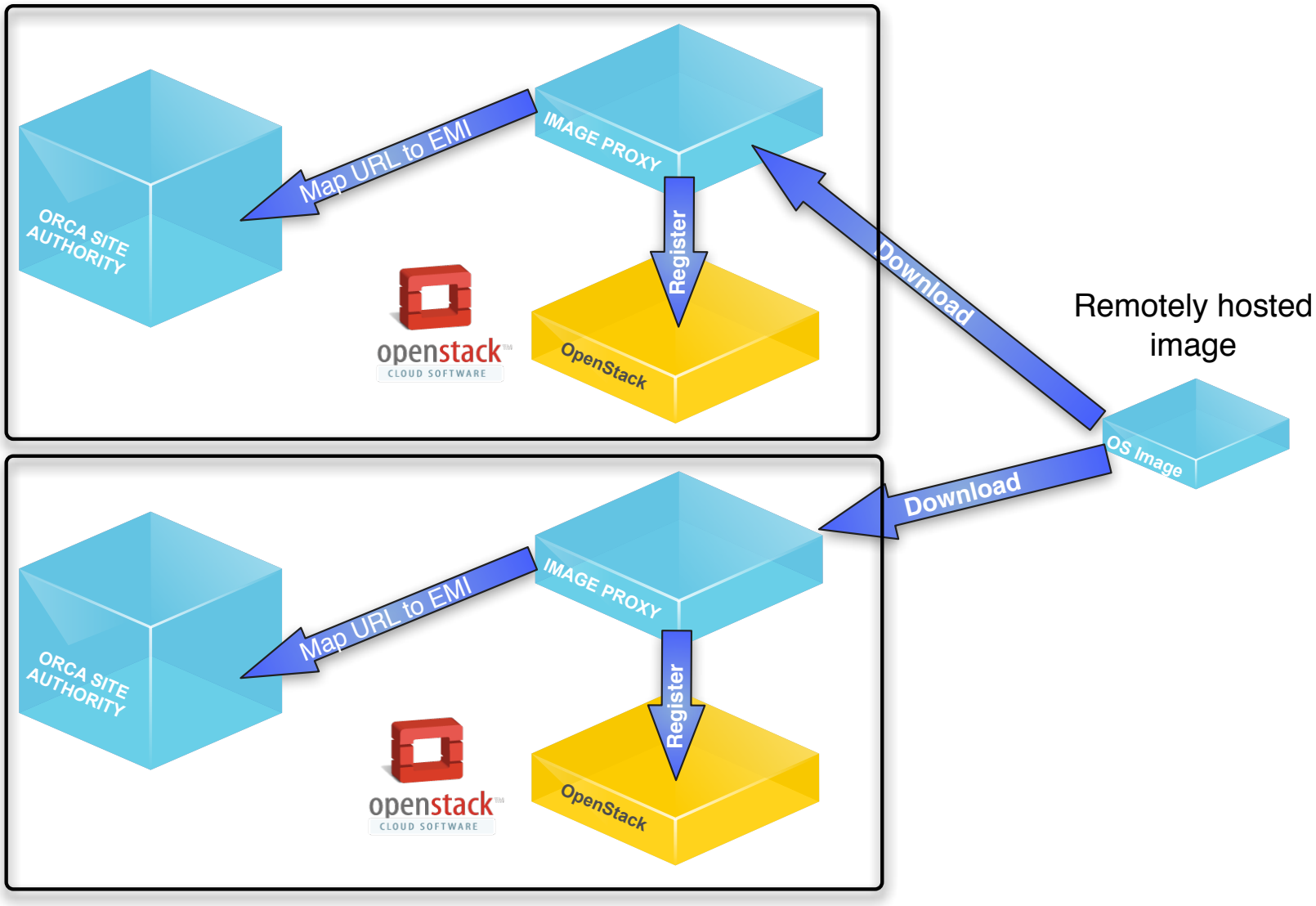
- Cut and paste slice name into the ‘Manifest View’ tab
- Click ‘Query for Manifest’
  - Inspect raw output if interested
  - Inspect the state of slice elements by right clicking on each element (usually ‘Ticketed’)
  - If you see ‘Failed’ you have a problem
- Poll by clicking ‘Query for Manifest’
  - Topology should materialize
  - All states should report ‘Active’
- Play around with layouts to get something pleasing



VM images

- Creating your own image
- Specifying your own image for ORCA
- Delays
  - Images are downloaded and registered with the site at the time of slice creation
    - If you repeatedly use the same image and the site already has it, this step is skipped
    - Images may be cached-out causing longer delays (to download and re-register)
- Are there examples of known good images?
  - Yes, visit https://geni-orca.renci.org/trac/wiki/neuca-images

VM Images


ORCA SITE AUTHORITY
Map URL to EMI
IMAGE PROXY
Register
openstack
CLOUD SOFTWARE
OpenStack
Download
Remotely hosted image
OS Image
Download
ORCA SITE AUTHORITY
Map URL to EMI
IMAGE PROXY
Register
openstack
CLOUD SOFTWARE
OpenStack

Example image metafile

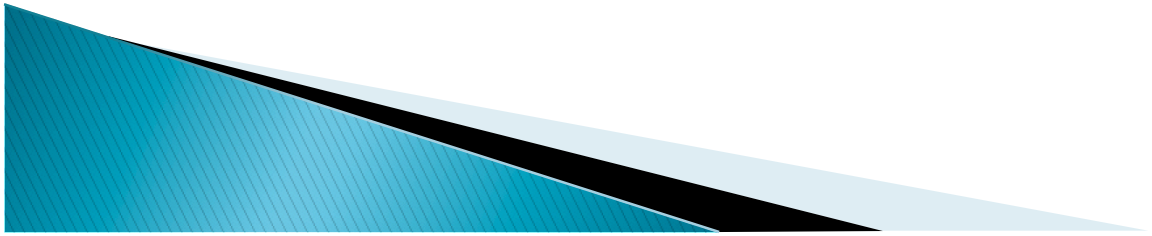
```
<images>
    <image>
        <type>ZFILESYSTEM</type>
        <signature>b54ed5a42cd99475c3d5d7c7a9839b69cf2076d5</signature>
        <url>http://geni-images.renci.org/images/workflows/pegasus/images/
    pegasus-4.0-v0.3.sparse.img.tgz</url>
    </image>
    <image>
        <type>KERNEL</type>
        <signature>f8a64d3bc429e8fb46c94ff3b11a932a27c142bc</signature>
        <url>http://geni-images.renci.org/images/workflows/debian-squeeze-
    kernel/vmlinuz-2.6.28-11-generic</url>
    </image>
    <image>
      <type>RAMDISK</type>
      <signature>6225968f43299aa40f6b1491360f3ce080bd16c4</signature>
      <url>http://geni-images.renci.org/images/workflows/debian-squeeze-kernel/
    initrd.img-2.6.28-11-generic</url>
    </image>
</images>
```

How does ORCA refer to an image

- URL of a metafile (can be same or different webserver as the image)
- SHA1 checksum of the metafile (to ensure it has not been modified)
- Workflow
  - Create filesystem, kernel ramdisk
  - Place on webserver
  - Take SHA1 signatures of each file
  - Generate metafile
  - Take SHA1 signature of metafile and its URL and add it to .flukes.properties or put it in Rspec
  - Try on a small slice (one node) to make sure it boots
- Can I put my image on your server?
  - Sorry, no.
- Will my image always remain cached at the racks?
  - No, depending on the use, your image may be cached out.

Doman binding

- Leaving domain as ‘System Select’ leaves ORCA to pick the domain that has available resources
- You can explicitly bind to specific domains
- If nodes or groups in slice request belong to different domains, appropriate inter-domain links will be provisioned on demand



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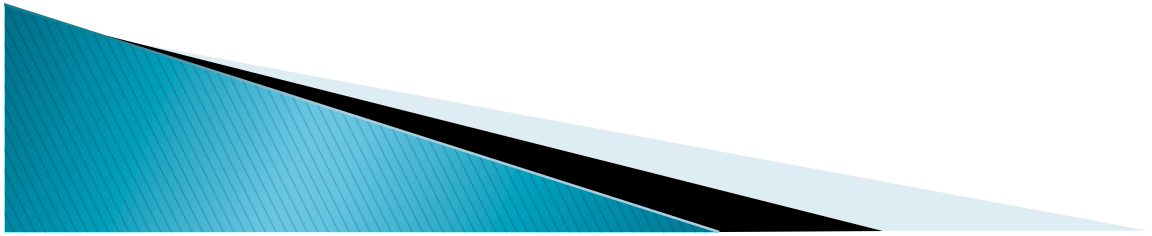
- Cut and paste slice name into the ‘Manifest View’ tab
- Click ‘Query for Manifest’
  - Inspect raw output if interested
  - Inspect the state of slice elements by right clicking on each element (usually ‘Ticketed’)
  - If you see ‘Failed’ you have a problem
- Poll by clicking ‘Query for Manifest’
  - Topology should materialize
  - All states should report ‘Active’
- Play around with layouts to get something pleasing

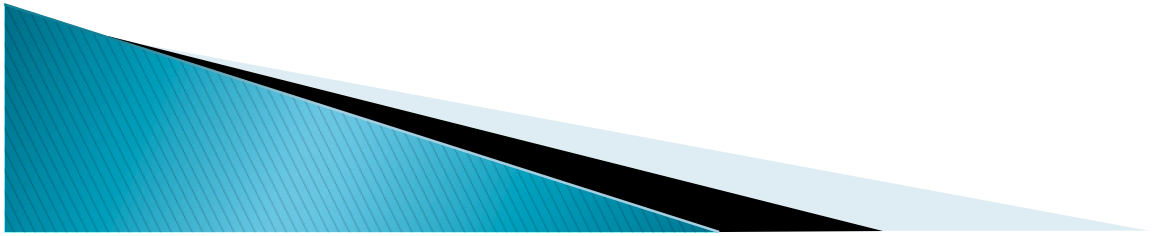


Logging into nodes

- Right click on node
- Select ‘Login to node’
  - In terminal window type in SSH key password (‘gec13’)
- Inspect uptime

  $ uptime
- Inspect the output of your boot script
- Inspect interfaces

  $ ifconfig
- Try to ping node neighbors



NEuca-py tools

- NEuca tools are loaded in the image
  - They configure network interfaces at boot time
  - They execute the post boot script
  - An image with NEuca tools will do neither of those things
    - You can still configure interfaces manually
- Allow you to inspect the VM configuration
- Run 'neuca' to get the list of neuca tools

  $ neuca
- Run 'neuca-user-data'
  - Note your boot script
- If you create your own VM image you are strongly encouraged to install NEuca tools on it
  - Visit https://geni-orca.renci.org/trac/wiki/NEuca-guest-configuration for instructions

Building your own image

- Build one from scratch using instructions from OpenStack or Eucalyptus
  - Virtio support is required
  - NEuca-py tools should be installed
- Build one by adding packages to one of the existing images
- Use post-boot scripts to install packages after the VM boots
  - E.g. wget a tar file or a deb or an RPM and install it
  - Beware some tools (e.g. apt-get) need a controlling TTY, which is not available when executed from post-boot script

Thank you for attending the tutorial!

- More Orca information
  - http://geni-orca.renci.org
- More ExoGENI information
  - http://wiki.exogeni.net