

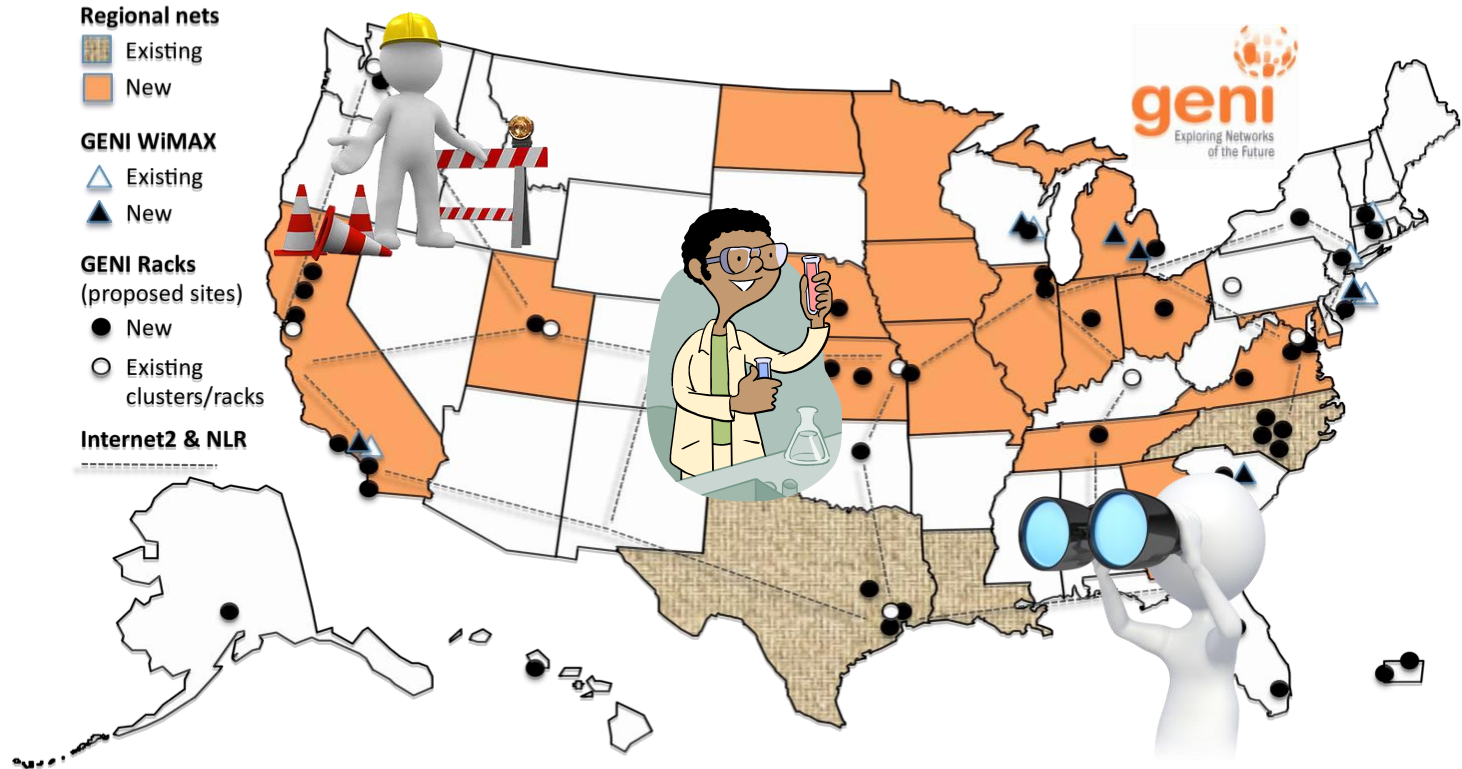
GENI

Train the TA – Session 1

Ben Newton, Jay Aikat, and Kevin Jeffay

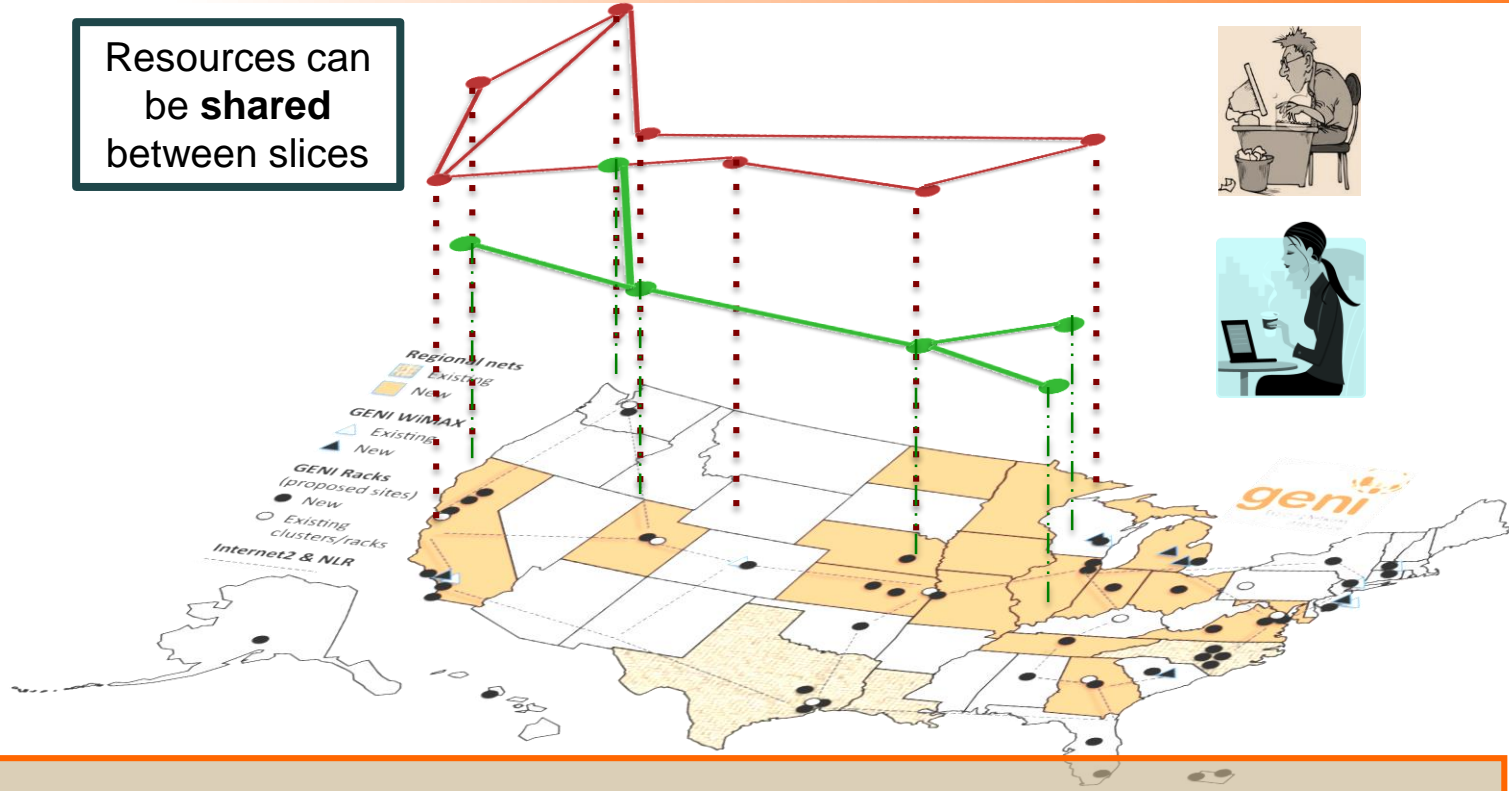
www.geni.net

- Welcome
- Introductions
- Tips for effective webinar
- Schedule
 - GENI Overview – Jay
 - GENI Terminology, SSH, Expiration and Renewal of Resources - Ben
 - Hands-on Lab – Ben and Jay
 - Behind the Scenes
 - Wrap-up



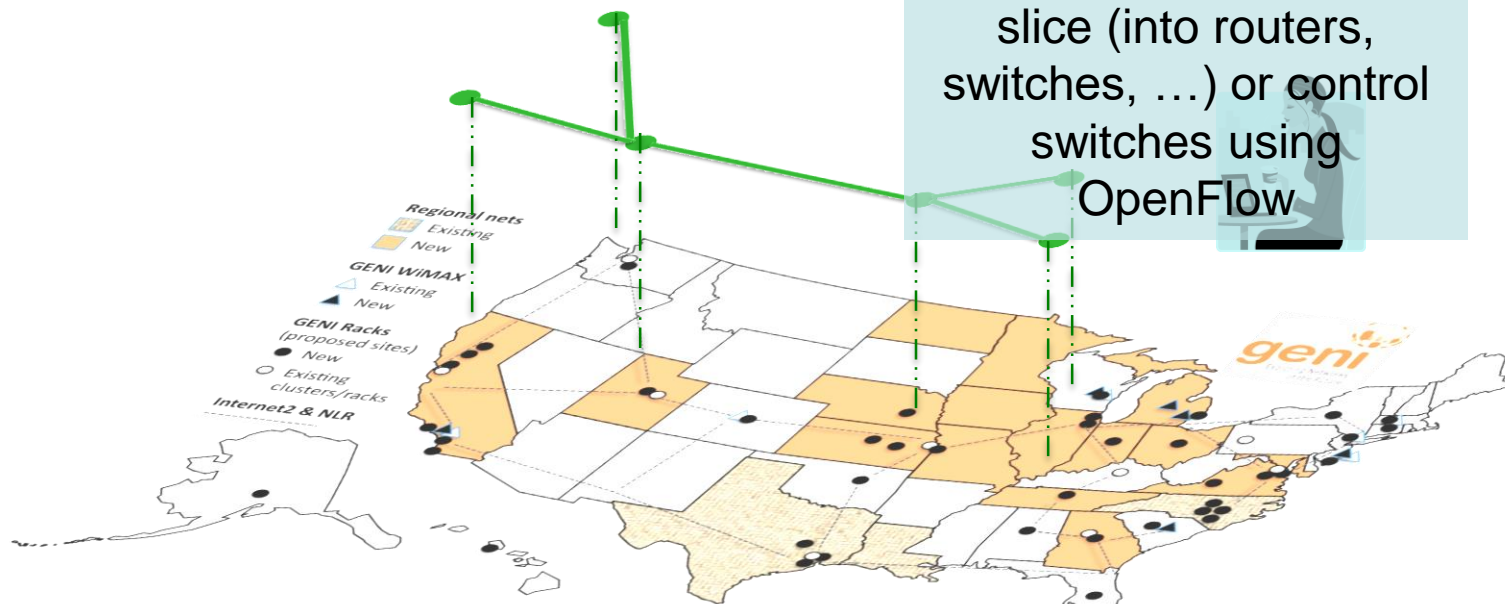
Multiple GENI Experiments run Concurrently

Resources can be **shared** between slices



Experiments live in **isolated** “slices”

I install software I want throughout my network slice (into routers, switches, ...) or control switches using OpenFlow



Experimenters can set up custom topologies, protocols and switching of flows

Compute Resources



GENI Racks: small clouds
Virtual Machines
Bare metal Machines



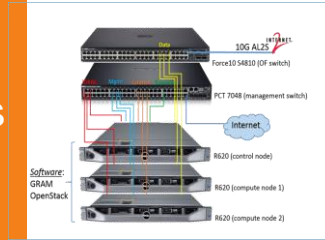
Android
Phones



Existing Testbeds

Network Resources

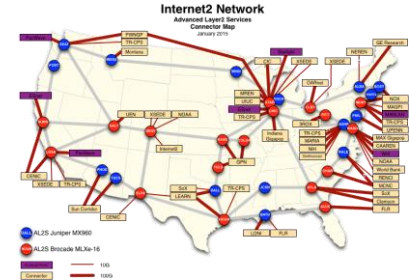
Layer 2 VLANs and Access to Programmable Switches



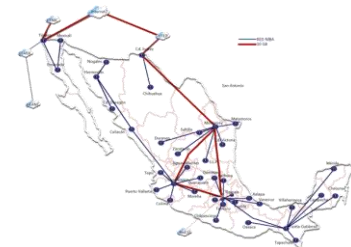
Rack switches



WiMAX/LTE
base stations,
4G/3G
Network



Internet2: US Research Backbone

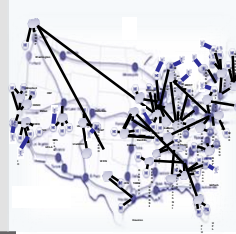


Regionals

GENI Use By Numbers



4300
Users



2600
Students
trained on
GENI by



200
+ Publications

45
Different
instructors



180
Universities

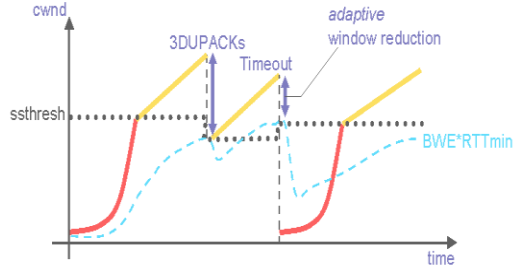


30
Countries



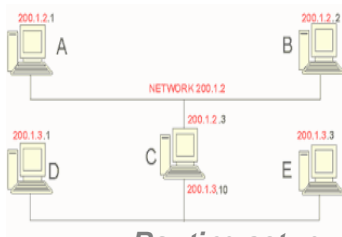
100,000+
Experiments
instantiated



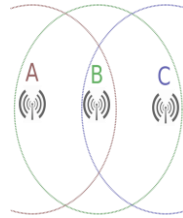


Congestion Avoidance (CA)
Slow-start (SS)

Protocol Behavior

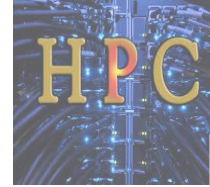


Routing setup



Wireless Communication

Teach Basic Concepts



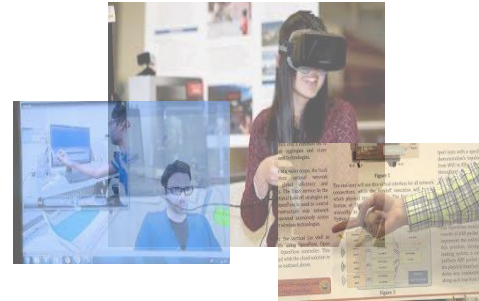
NFV

SDN

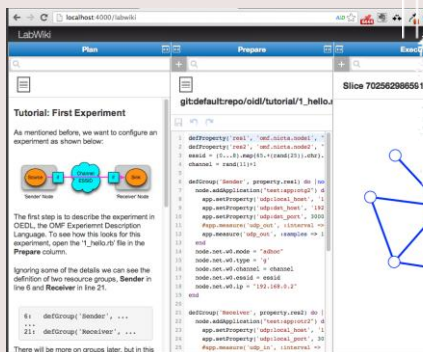
OpenFlow

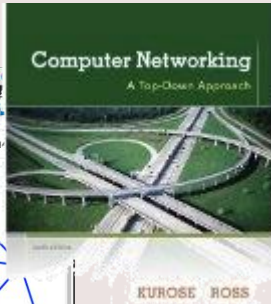


New technologies



Semester Projects





Labs on GENI for networking textbook

Mike Zink
UMass Amherst



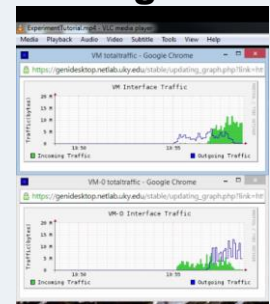
Massive Online Open Courses on GENI



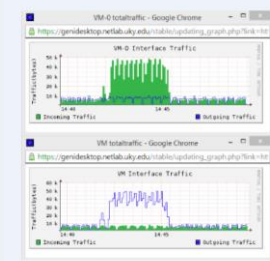
Shivendra Panwar,
Thanasis Korakis
NYU Poly

Use GENI to educate the Internet users, not the Internet creators.

GENI Modules to teach networking concepts



Example Demo Module



Example Assignment
Kevin Jeffay, Jay Aikat
UNC-Chapel Hill

GENI Terminology



Sponsored by the National Science Foundation

www.geni.net



slice

project

experimenter

resource

aggregate

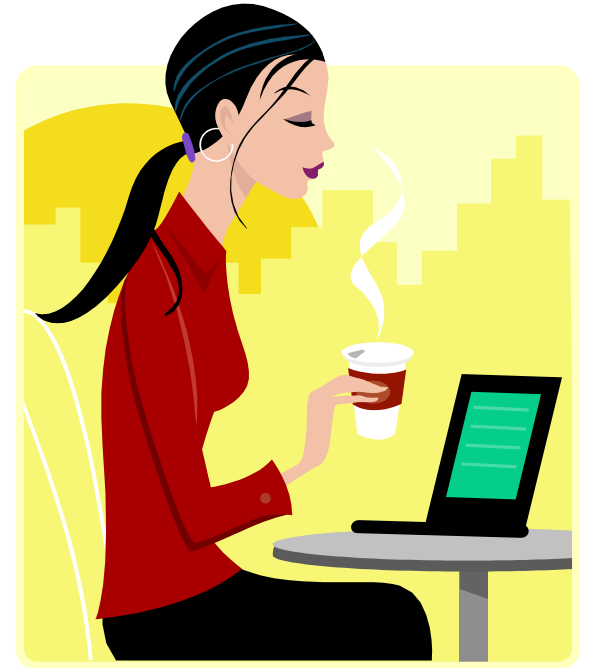


An **experimenter**

is a researcher who uses GENI resources

Different types of experimenters have different roles and permissions:

- Advisor vs Grad Student
- Teacher vs TA vs Student



Experimenter



Creating an Account

Use GENI

GENI Portal is at:

<https://portal.geni.net>



Sponsored by the National Science Foundation

www.geni.net



GENI User Authentication

The GENI Portal leverages InCommon for single sign-on authentication

InCommon[®]

Students from 518 educational and research institutions have InCommon accounts

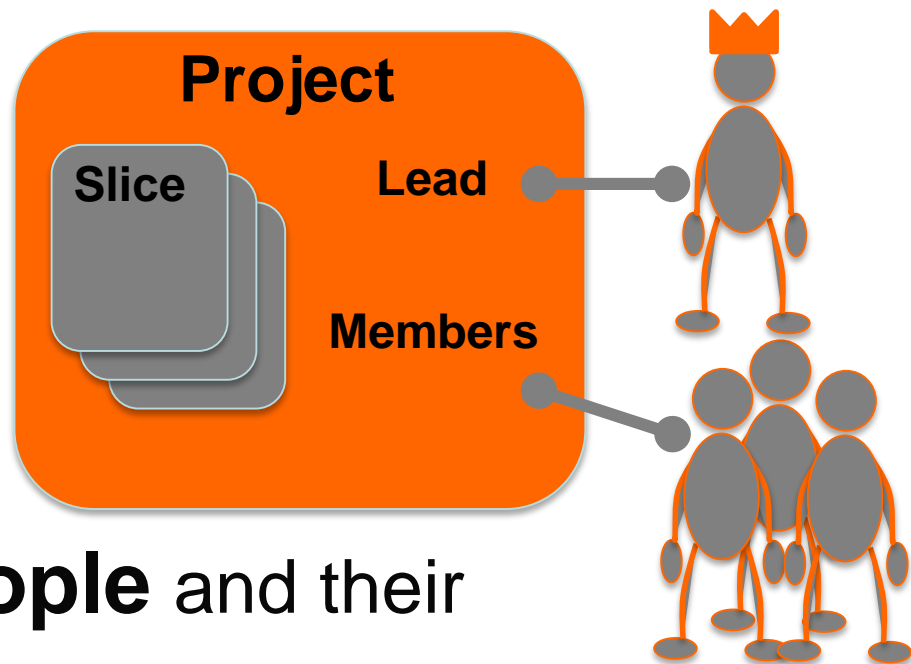
For many experimenters:

- no new passwords
- familiar login screens

GENI Project Office runs a federated IdP to provide accounts for non-federated organizations.



Projects organize research in GENI



Projects contain both **people** and their **experiments**

A project is led by a single responsible individual:
the **project lead**



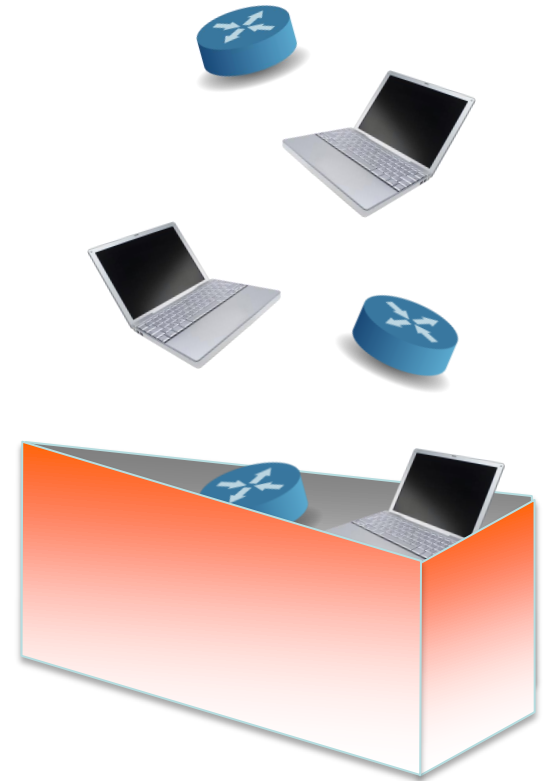
A **slice** is a *container* of resources used in an *experiment*.

A slice can contain resources from one or more aggregates

A slice is in a single project

A slice has an ***expiration***

Slice names are ***public, reusable*** and ***unique*** (*within a project*)



A resource

is a piece of infrastructure

A resource can be real or virtual.

Resource specifications (aka. **RSpecs**) are used to describe and request resources.

Examples:

- Compute: computer vs virtual machine (VM)
- Wireline Network: VLAN or OpenFlow
- Wireless: WiMAX



An **aggregate** manages a set of reservable **resources**

Aggregates include:

GENI racks

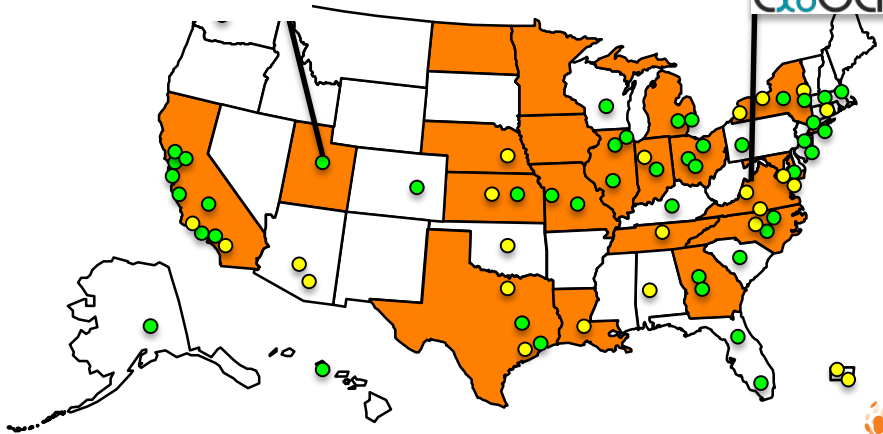
OpenFlow

WiMAX

InstaGENI Rack



ExoGENI Rack



project

Lead:



**Experimenter
(aka Professor)**

Member:



**Experimenter
(aka Student)**

slice

aggregate



How SSH *With a Private Key* Works



Sponsored by the National Science Foundation

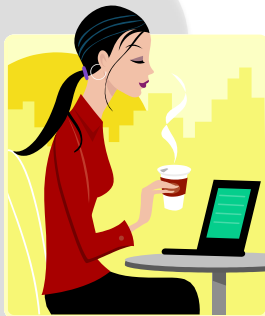
www.geni.net



SSH with a *password*

*nix-based system (Windows behavior may vary)

```
local> ssh jdoe@remote.edu
jdoe@remote.edu's password:
#####
Welcome to remote!
jdoe@remote> exit
local> ssh jdoe@remote2.edu
jdoe@remote2.edu's password:
#####
```



ssh



User enters password
once for
each connection to
each machine

Hash of password
stored on each
remote machine



SSH with a *private key*

*nix-based system (Windows behavior may vary)

```
local> ssh-add ~/.ssh/id_rsa
Enter passphrase for ~/.ssh/id_rsa:
#####
local> ssh jdoe@remote.edu
Welcome to remote!
jdoe@remote> exit
local> ssh jdoe@remote2.edu
Welcome to remote2!
jdoe@remote2> exit
local> ssh jdoe@remote3.edu
Welcome to remote3!
jdoe@remote3> exit
local> ssh-add -D
```



User enters passphrase to unlock private key for *all* connections to *all* machines

Private key is stored only on local machine

Public key is stored on each remote machine





SSH with a private key is both
easier to use and more secure



Sponsored by the National Science Foundation



Sponsored by the National Science Foundation



You should *never* be prompted for a password to log into a GENI compute node.

If you are, something has *always* gone wrong.



Sponsored by the National Science Foundation

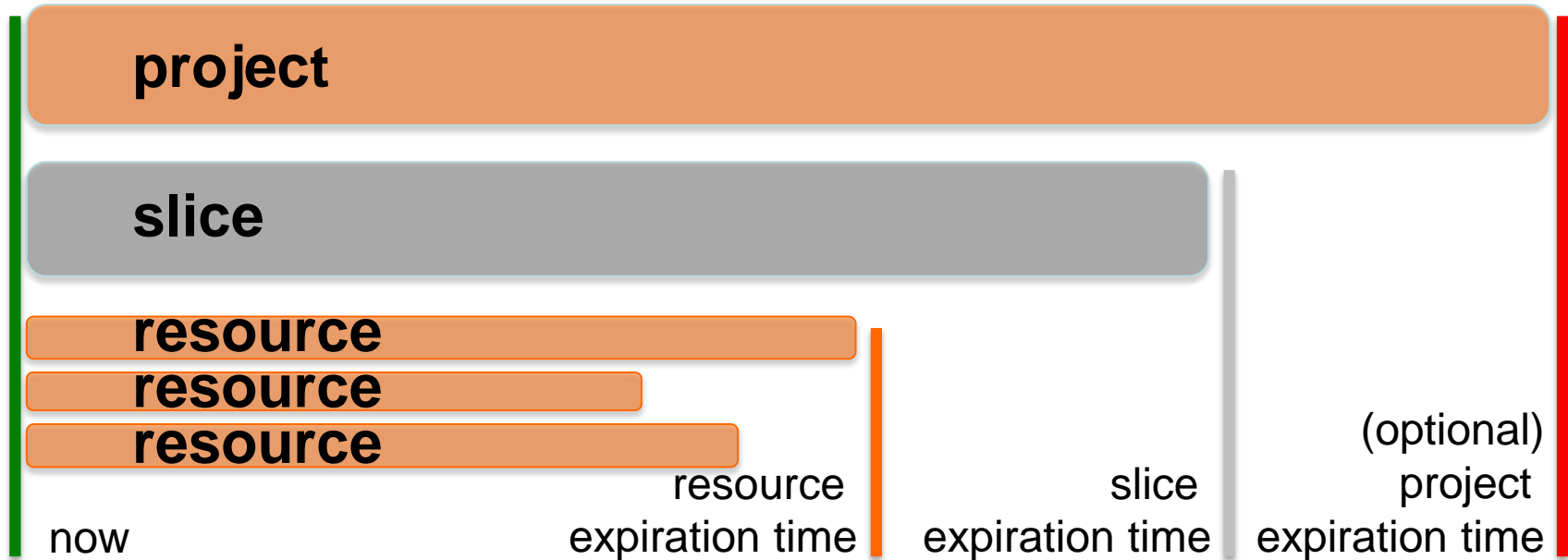


Sponsored by the National Science Foundation

GENI Expiration and Renewal



Sponsored by the National Science Foundation



To extend the lifetime of your resource reservation, you must renew the **slice** and **all resources**



Are you ready for the tutorial?

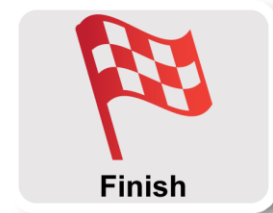
1. Grab a Worksheet and instructions
2. Did you do the pre-work?
 - A. Do you have an account?
 - B. Have you installed the tools?
 - * SSH
3. Connect to the network

GENI Portal is at:

<http://portal.geni.net>

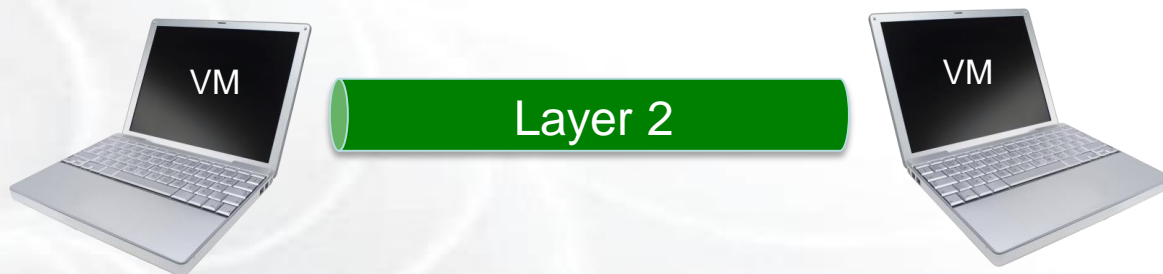
Lab Zero: A First Experiment using GENI

Ben Newton, Jay Aikat, and Kevin Jeffay
UNC Chapel Hill



Do a Simple Experiment in GENI

Reserve two VMs connected at Layer 2





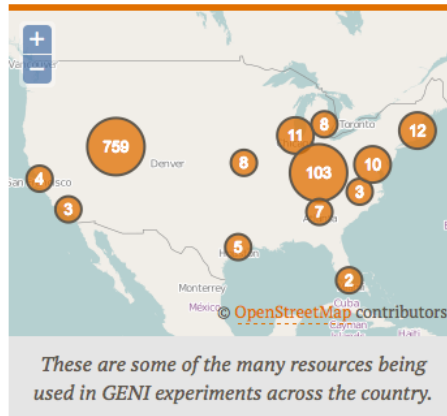
WELCOME TO GENI

GENI is a new, nationwide suite of infrastructure supporting "at scale" research in networking, distributed systems, security, and novel applications. It is supported by the [National Science Foundation](#), and available without charge for research and classroom use.

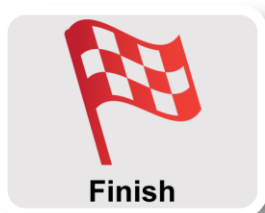
Use GENI

Find out more about using GENI

- [Information for GENI experimenters](#)
- [Published research that used GENI resources](#)
- [Get help using GENI](#)



Experiment Workflow



- Part I: Design/Setup

- Part II: Execute

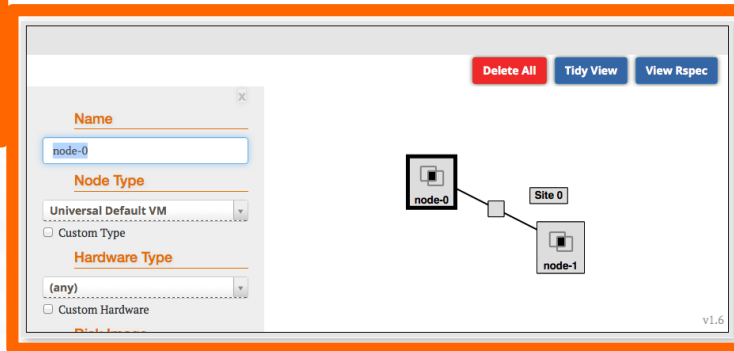
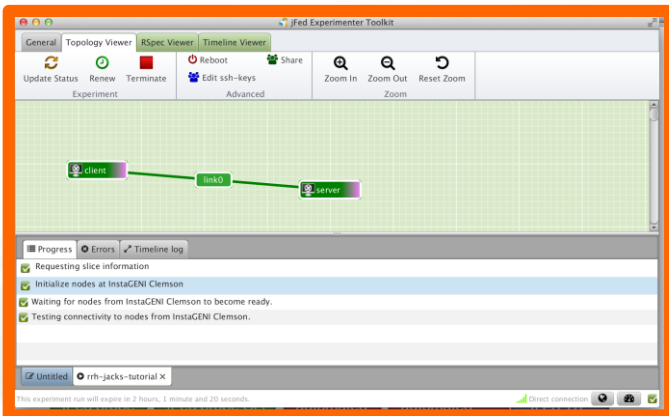
- Part III: Finish

The GENI Portal is...

Use GENI

A web-based tool for experimenters to manage
experimenters, projects, and slices.

Includes simple tools to reserve **resources.**

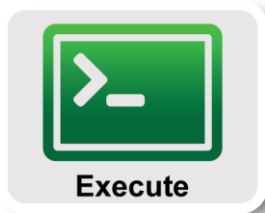


Graphical user interfaces (GUIs) for:

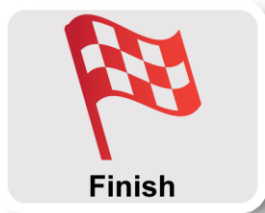
- **designing topologies** in GENI
- **reserving resources** in GENI



- **Part I: Design/Setup**



- **Part II: Execute**



- **Part III: Finish**

Establish Management Environment

Use GENI

1 Pre-work: Design your experiment

2.1 Pre-work: Create a GENI account

2.2 Pre-work: Project lead (aka professor) adds you to project

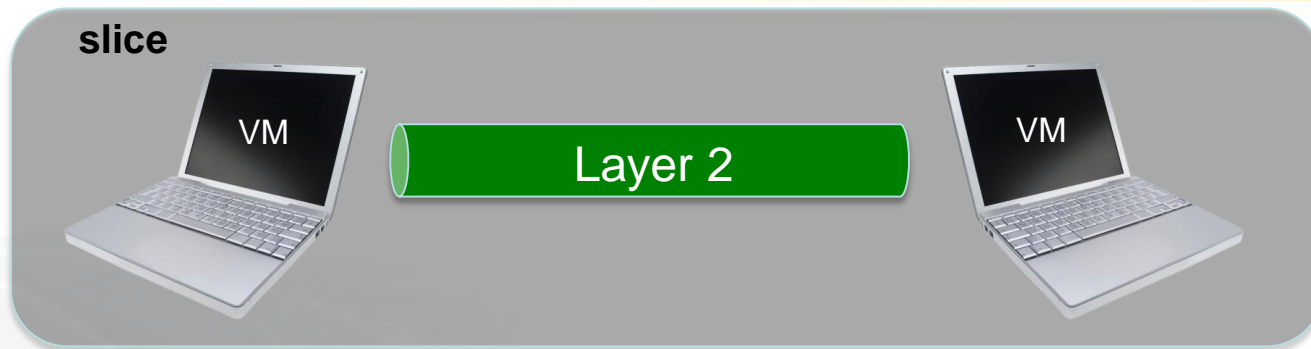
Project Name: TrainTheTA

2.3 Generate and Download SSH Keypair

2

On your local machine...

```
> mv ~/Downloads/id_geni_ssh_rsa  
~/ssh/.  
> chmod 600 ~/ssh/id_geni_ssh_rsa  
> ssh-add ~/ssh/id_geni_ssh_rsa
```

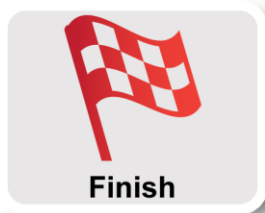


3.1 Create a slice

3.2 (optional) Renew your slice

3.3 Reserve two VMs at one aggregate

3.4 Check Whether VMs are Ready to be Used

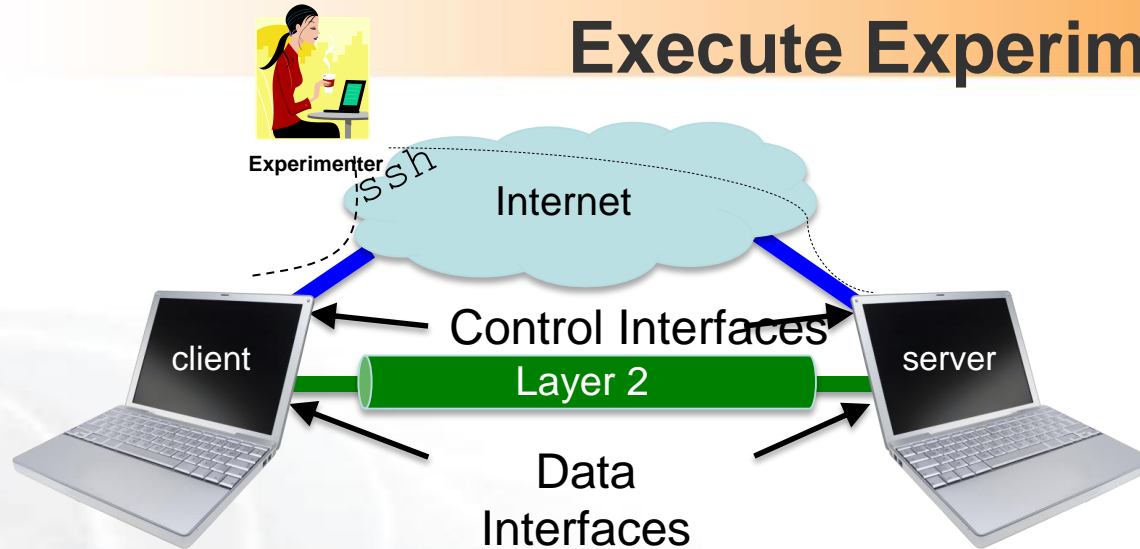


- Part I: Design/Setup

- **Part II: Execute**

- Part III: Finish

Execute Experiment



4.1 Login to all three nodes

5.1 Test Connectivity

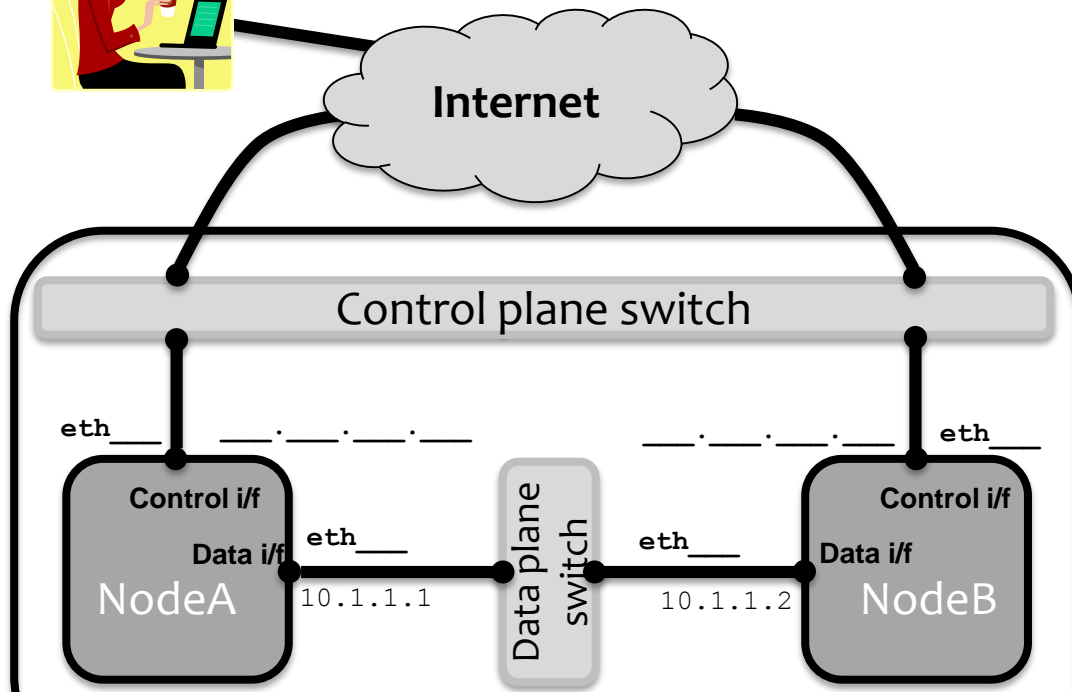
5.2 Explore the Data and Control Planes

6.1 Logout of nodes

Worksheet

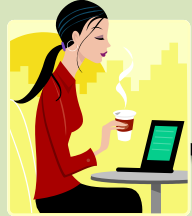
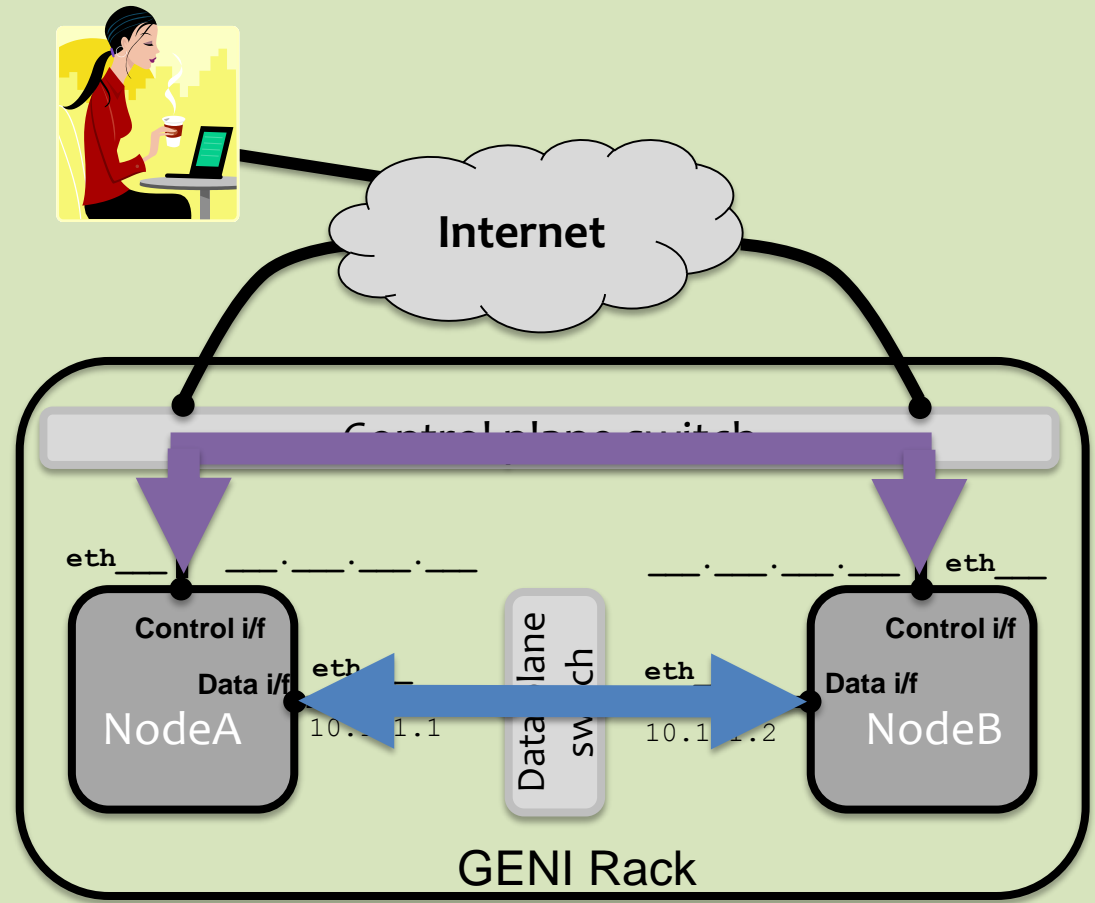
Slice Name:

lab0<your initials>

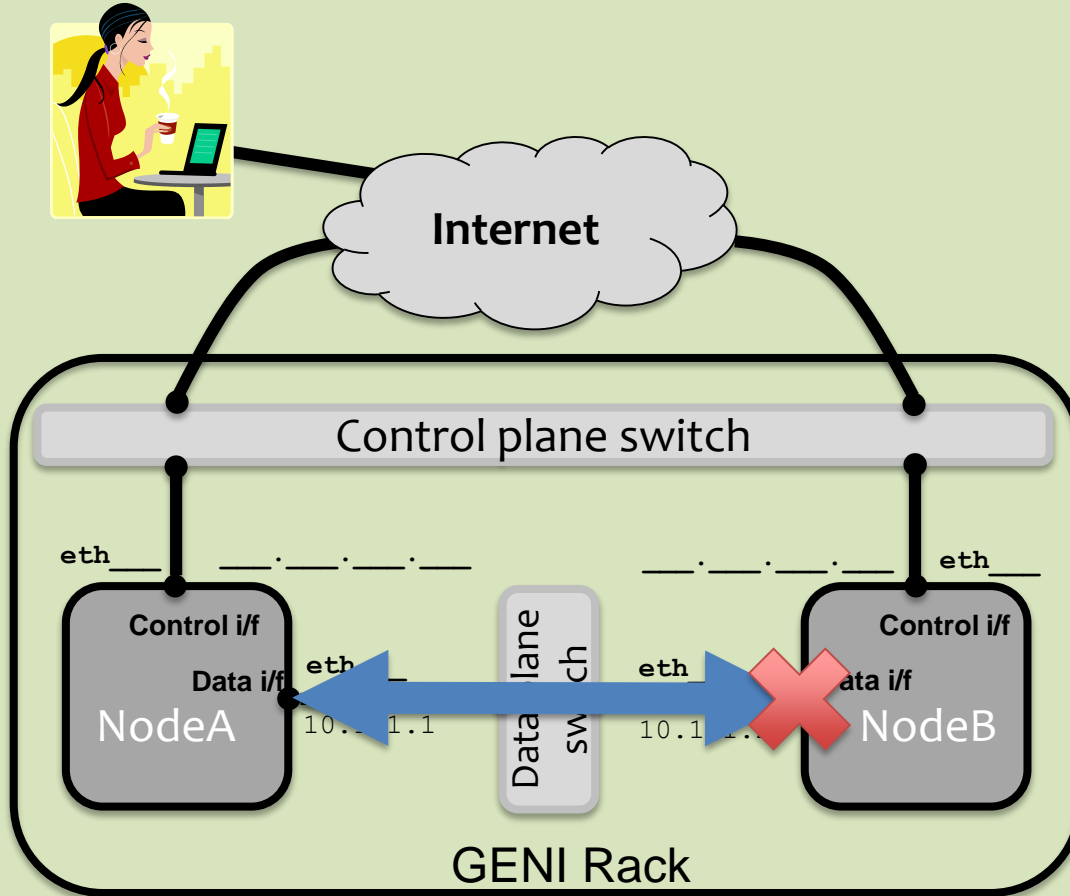


5.1

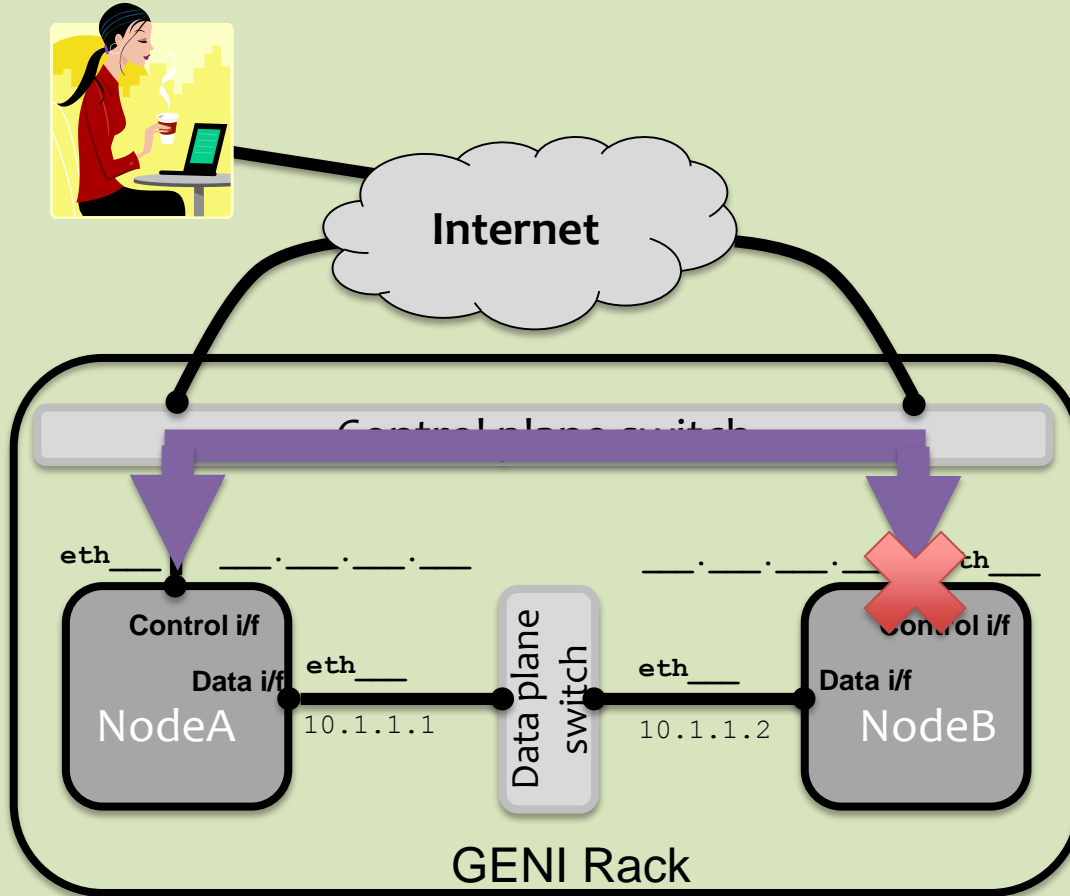
What is the bandwidth of the **data** link? Why?
What is the bandwidth of the **control** link? Why?



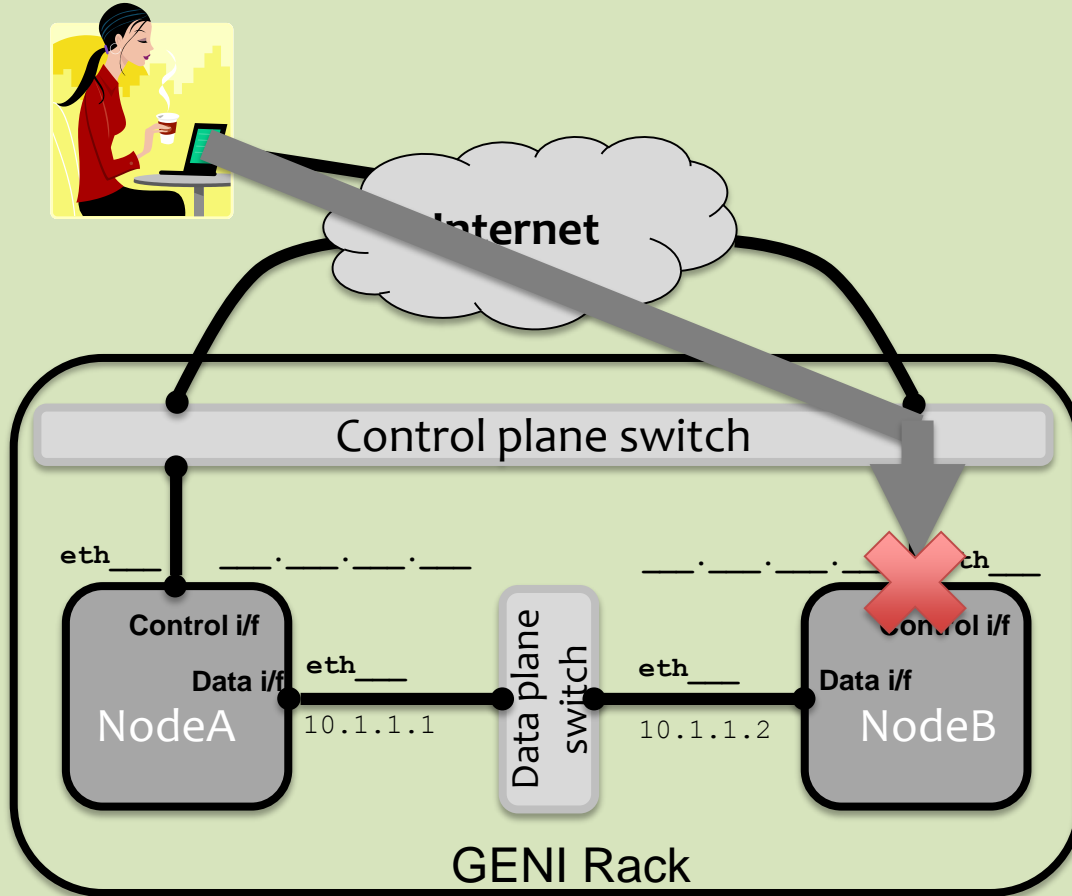
5.3 When you bring down the **data** interface, the destination should become unreachable. Why?



- 5.3 After you bring down the **control** interface, the destination becomes unreachable. Why?

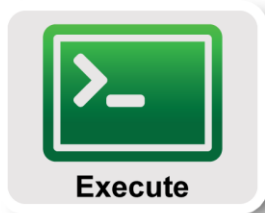


- 5.3 After you bring down the **control** interface, your ssh session should immediately hang. Why?





- Part I: Design/Setup



- Part II: Execute



- Part III: Finish

7

JFed Experimenter Toolkit

General Topology Viewer RSpec Viewer Timeline Viewer

New Open Open URL Save Run Update Status Terminate Recover Preferences Report a bug Documentation

Experiment Definition Experiment Preferences Support

client link0 server

Progress Errors Timeline log

- Requesting slice information
- Initialize nodes at InstaGENI Clemson
- Waiting for nodes from InstaGENI Clemson to become ready.
- Testing connectivity to nodes from InstaGENI Clemson.

Untitled rrh-jacks-tutorial x

This experiment run will expire in 1 hour, 58 minutes and 45 seconds.

Delete Resources

Flack GENI Desktop LabWiki Omni jFed

Current Resources Resource Details

Manage Resources

Resources on RENCI ExoGENI are ready.

server client

RENCI ExoGENI

Delete known slice resources?

Cancel OK

Renew Renew Date Delete SSH Restart Details Status Add Resources

project resource
aggregate experimenter



When your experiment is done, you should always release your resources.

- Normally this is when you would archive your data
- Delete your resources at **each** aggregate

You have...

- Run your first GENI Experiment!
- Exercised your knowledge of GENI terminology
- Used the GENI Portal and Jacks or jFed



Welcome to GENI!

Behind the Scenes of GENI Experimentation

Understanding GENI AM API and RSpec



Sponsored by the National Science Foundation

What happens behind the scenes when reserving resources on GENI?



All GENI resource reservation tools use the same data formats and APIs

RSpecs

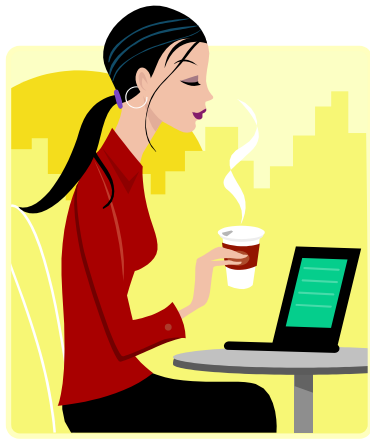
GENI Resource Specifications

GENI AM API

GENI Aggregate Manager API



Sponsored by the National Science Foundation



researcher tool



create slice

CH APIs

slice credential

slice, RSpec

Aggregate Manager API



clearinghouse



aggregate resources



GENI Resource Specifications (or Rspecs) are XML documents that describe resources

```
<?xml version="1.0" encoding="UTF-8"?>
<rspec type="request" ...
  xmlns="http://www.geni.net/resources/rspec/3">
  <node client_id="server"
component_manager_id="urn:publicid:IDN+instageni.gp
olab.bbn.com+authority+cm">
    <sliver_type name="emulab-xen"/>
    <interface client_id="server:if0"/>
  </node>
</rspec>
```

RSpec for a virtual machine with one interface



Three flavors of RSpec

Advertisement RSpecs

Aggregates describe what they have

Request RSpecs

Experimenters describe the resources they want

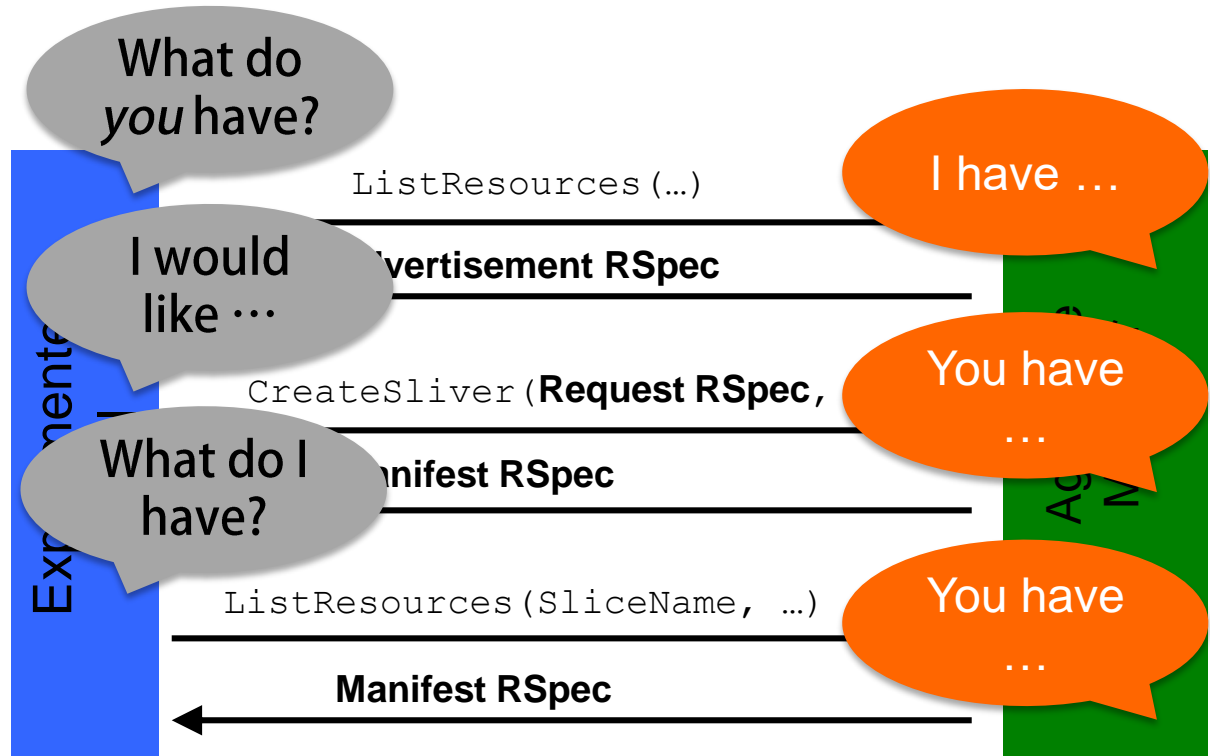
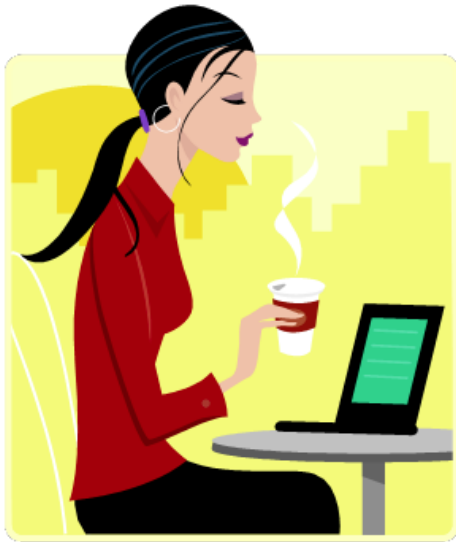
Manifest RSpecs

Aggregates describe resources allocated to an experimenter



Experimenter tools and aggregates talk to each other using the GENI Aggregate Manager API or

GENI AM API





All GENI resource reservation tools use the same data format (RSpec) and APIs (AM API)

Pick the tool that works best for the current task

An Introduction to GENI Tools

Jacks in the Portal and omni



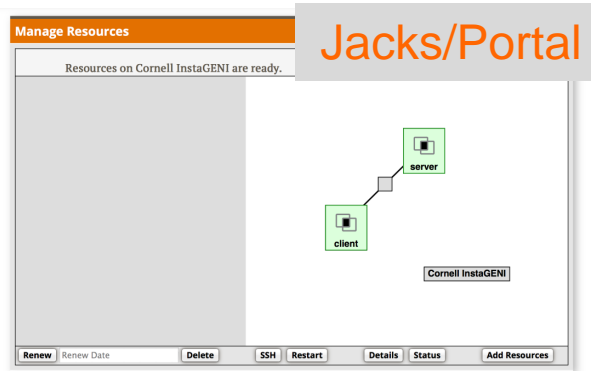
Sponsored by the National Science Foundation

www.geni.net



All GENI resource reservation tools speak AM API and RSpec...

Hosted



The screenshot shows the 'Manage Resources' page in the Jacks/Portal interface. The title 'Jacks/Portal' is overlaid in orange. The main content area displays a network diagram with a 'client' node connected to a 'server' node, both represented by server icons. Below the diagram, the text 'Resources on Cornell InstaGENI are ready.' is visible. At the bottom, there is a control bar with buttons for 'Renew', 'Delete', 'SSH', 'Restart', 'Details', 'Status', and 'Add Resources'.

Local

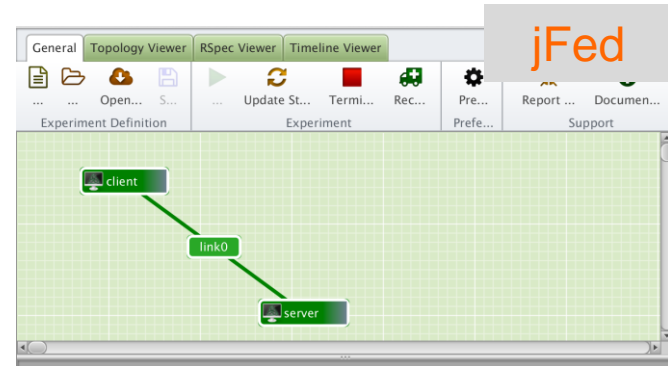
```
$ omni createsliver aliceslice myRS
INFO:omni: -----
INFO:omni: Completed createsliver:

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
INFO:omni: =====
```

omni



The screenshot shows the jFed web interface. The title 'jFed' is overlaid in orange. The interface includes a menu bar with options like 'General', 'Topology Viewer', 'RSpec Viewer', and 'Timeline Viewer'. Below the menu, there are icons for file operations and experiment management. The main area displays a network diagram with a 'client' node connected to a 'server' node via a 'link0' link, all on a green grid background.



GENI Tool: Jacks in the Portal

A graphical user interface

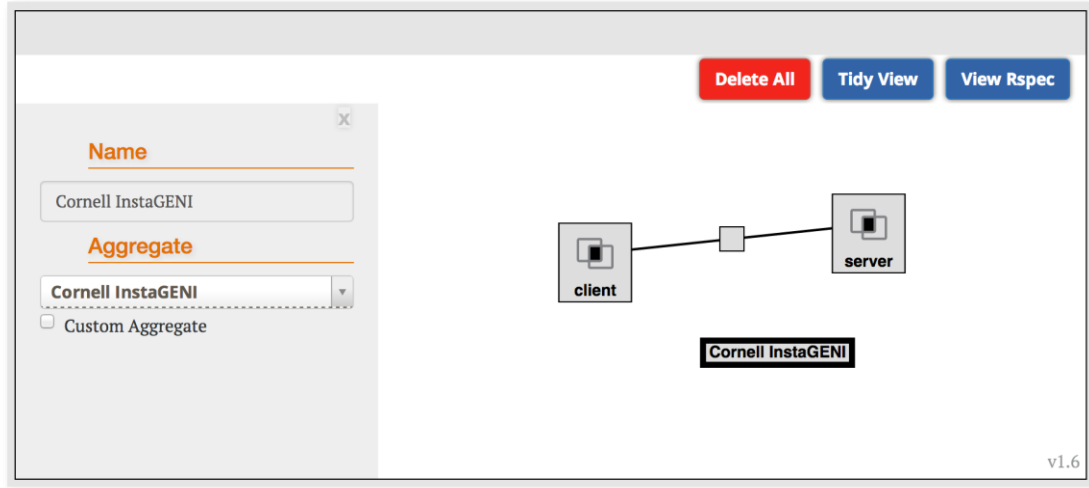


Sponsored by the National Science Foundation

www.geni.net



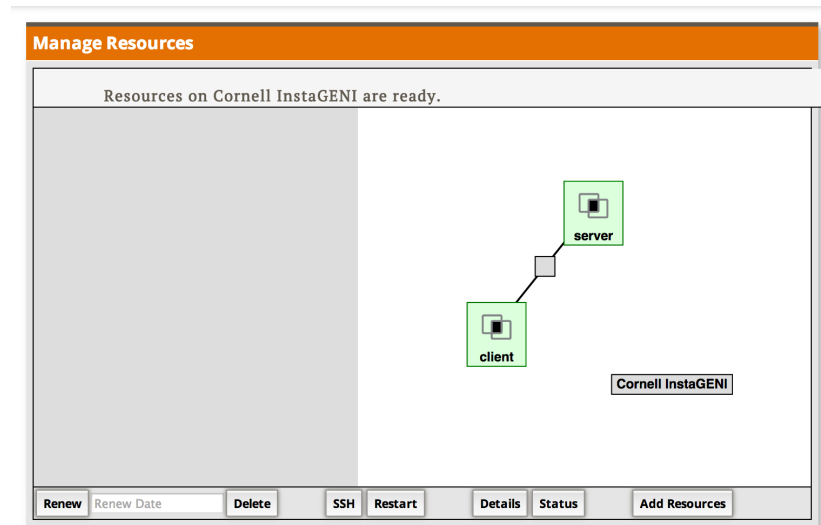
When you draw a topology using a tool, it creates a **request RSpec** describing the resources

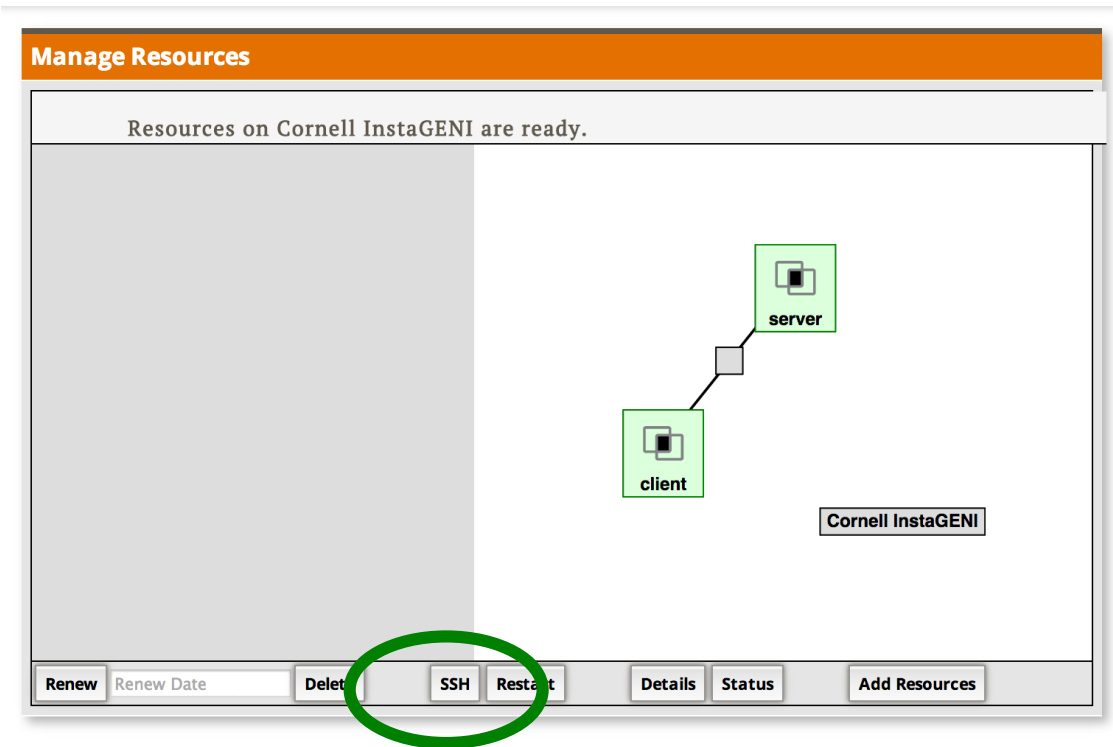


When you reserve resources at an aggregate, **createSliver** is called using that request

Tools periodically call `sliverStatus` at each used aggregate

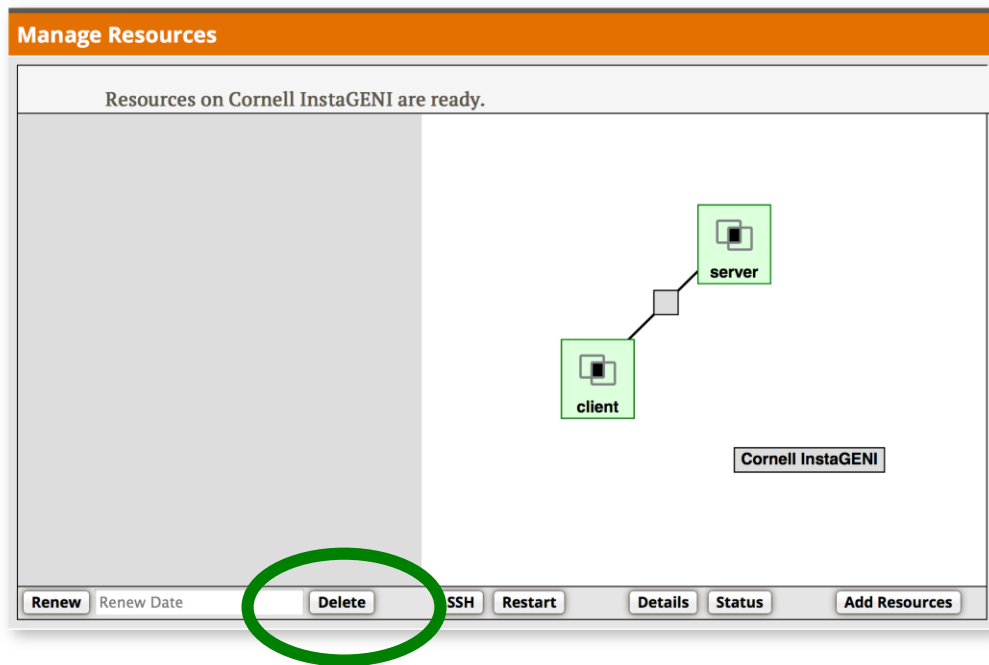
When the returned status is “ready”, the resources turn green





ListResources on a slice
returns a **manifest RSpec**

Manifest includes info to
log into VMs



When you delete resources, tool calls `deleteSliver` on aggregates

GENI Tool: Omni

A command line tool



Sponsored by the National Science Foundation

omni Resource Reservation Tool

A command line tool for making AM API calls at *any* aggregate that implements the GENI AM API

Written in and scriptable from Python

```
$ 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
```

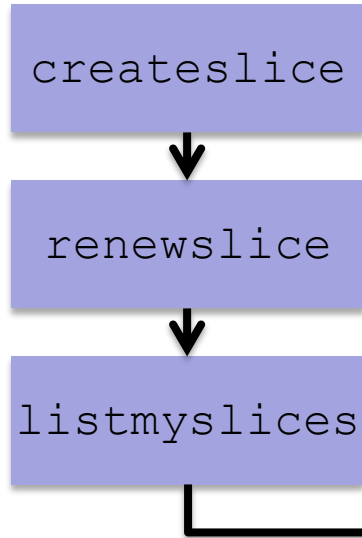
Args: createsliver aliceslice myRSpec.xml

```
Result Summary: Slice urn:publicid:IDN+pgeni
Reserved resources on https://www.emulab.net/p
Saved createsliver results to aliceslice-man
INFO:omni: =====
```



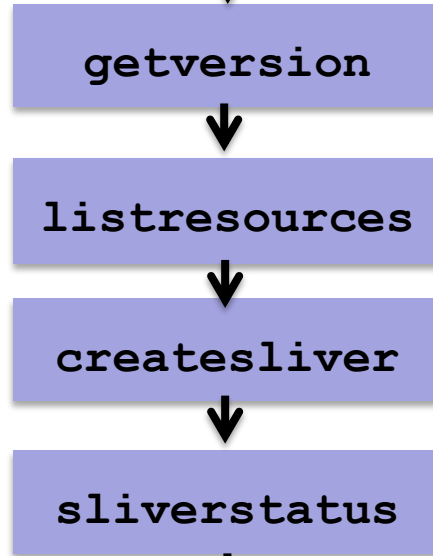
omni Workflow

Create Slice



Create Sliver

Repeat for each aggregate



Cleanup

Repeat for each aggregate



Sample Commands

```
omni -a aggregatename listresources
```

```
omni -a aggregatename createsliver slicename requestRSpec
```

```
omni -a aggregatename sliverstatus slicename
```

```
omni -a aggregatename listresources slicename
```

```
omni -a aggregatename deletesliver slicename
```

```
readyToLogin -a aggregatename slicename
```

Returns the ssh commands needed to log into nodes



Mentioned in this talk

GENI Portal

`https://portal.geni.net`

Omni

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



Sponsored by the National Science Foundation

GENI Support for Automating Resource Reservation



Sponsored by the National Science Foundation

There are three basic techniques for automating experiment configuration:

- OS images
- Install scripts
- Configuration management tools



Operating System Images

use standard images as is

OR

snapshot them to create custom images

```
<node client_id="node" ...>  
  <sliver_type name="emulab-xen">  
    <disk_image name="...+image+emulab-ops:UBUNTU12-64-STD"/>  
  </sliver_type>  
</node>
```





DO use custom OS images if the software installation is complicated or slow



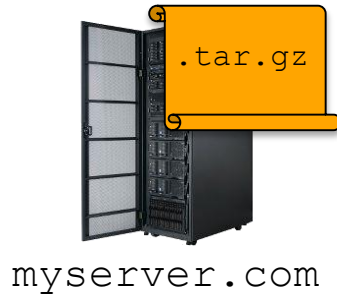
DO NOT use custom OS images as a substitute to being able to reproduce the installation



Install and Execute Scripts



Install and Execute Scripts



```
<node client_id="node" ...>
```

```
<services>
```

```
<install url="http://myserver.com/mycode.tar.gz"  
install_path="/local" />
```

```
<execute command="sudo /local/install-script.sh".../>
```

```
</services>
```

```
</node>
```





DO use install scripts for frequently used configurations which must run at start time



DO NOT use install scripts for infrequent tasks



Configuration Management (CM) Tools

- Ensure experiment is in a known configuration
- Easily reproduce experiment configurations for multiple runs, changing parameters, scaling up, etc

Ansible and Chef open source CM tools are regularly used with GENI





DO use configuration management tools for every day tasks and to ensure repeatable experiments





Automate your experiments using a combination of

- *OS image,*
- *install scripts, and*
- *configuration management tools*



Sponsored by the National Science Foundation

www.geni.net



Next TA-Training Webinar @ 2:00 PM on Fri, Feb 12

In the meantime, please try the Take-home assignment on:
<http://groups.geni.net/geni/wiki/GENIExperimenter/Tutorials/TrainTheTA-Spring2016>

If you have questions, please contact:
Ben Newton (bn@cs.unc.edu)