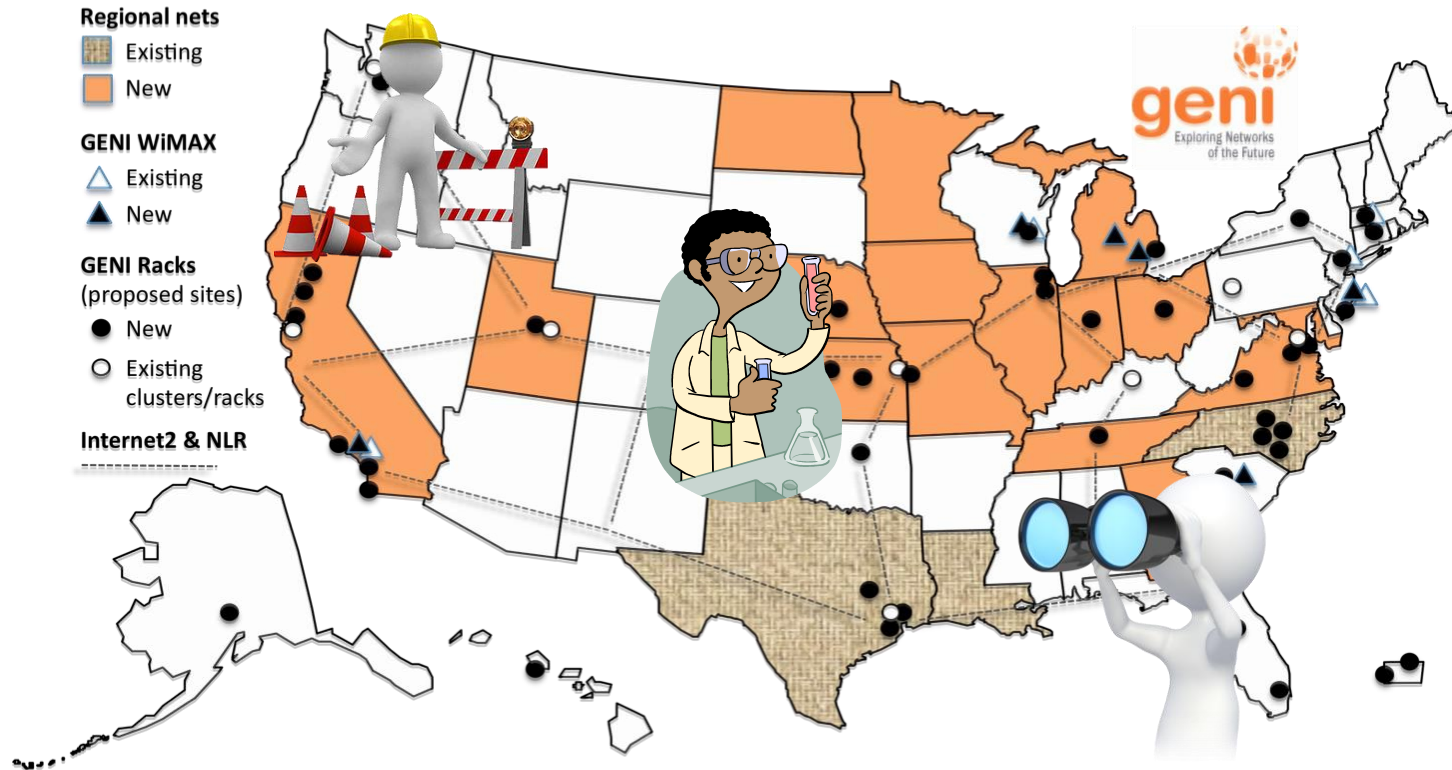


# GENI in the Classroom – Session 1

**Ben Newton, Jay Aikat, and Kevin Jeffay**

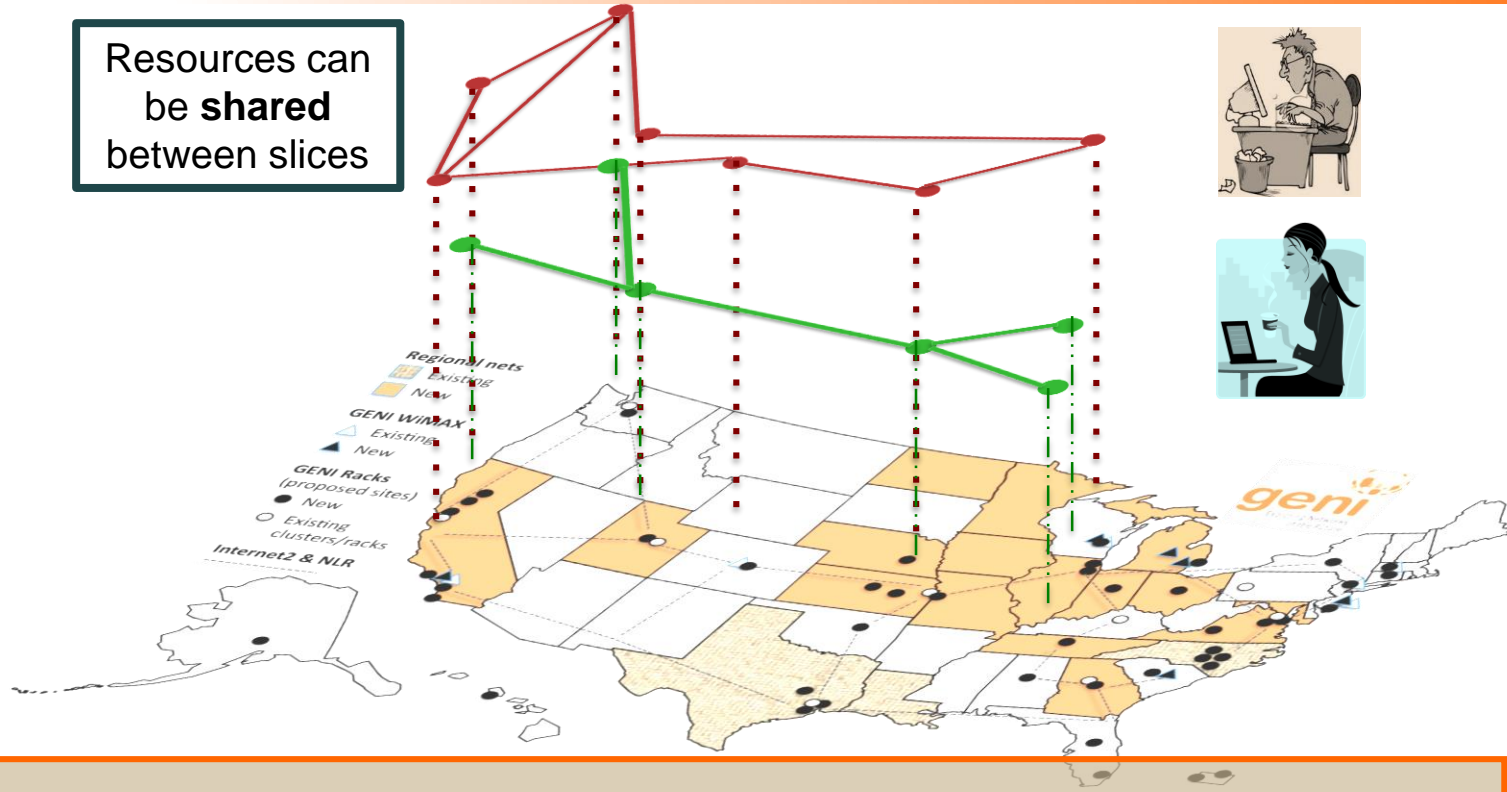
**[www.geni.net](http://www.geni.net)**

- 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 - Ben
  - Wrap-up



# Multiple GENI Experiments run Concurrently

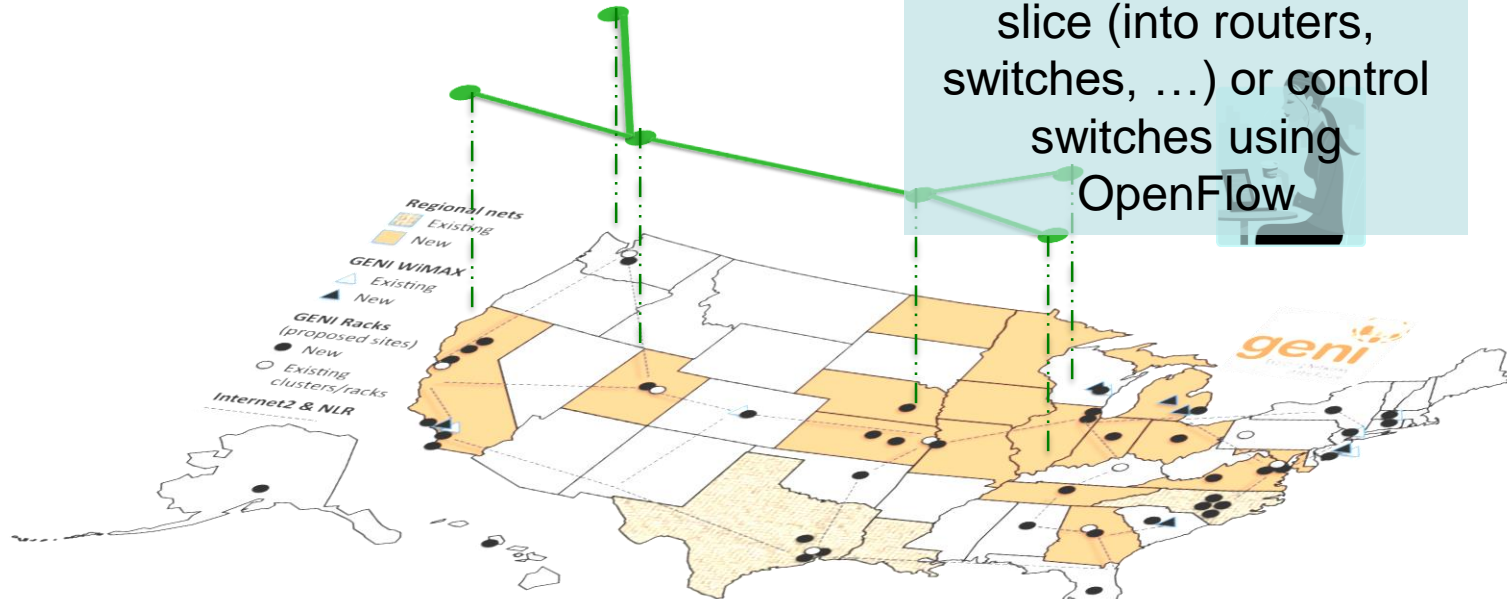
Resources can  
be **shared**  
between slices



Experiments live in **isolated** "slices"

# GENI is “Deeply Programmable”

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



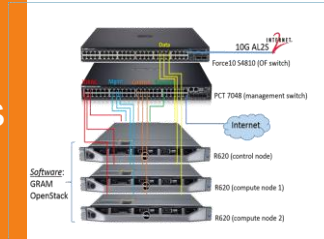
Android  
Phones



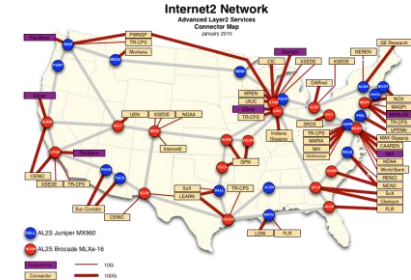
Existing Testbeds

# Network Resources

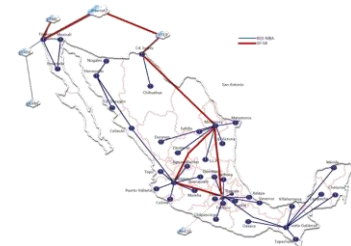
Layer 2 VLANs and Access to Programmable Switches



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



Internet2: US Research Backbone



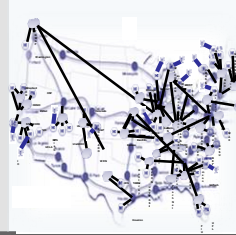
Regionals



# GENI Use By Numbers



**4300**  
Users



**2600**  
Students  
trained on  
GENI by



**200**  
+  
Publications

**45**  
Different  
instructors



**180**  
Universities

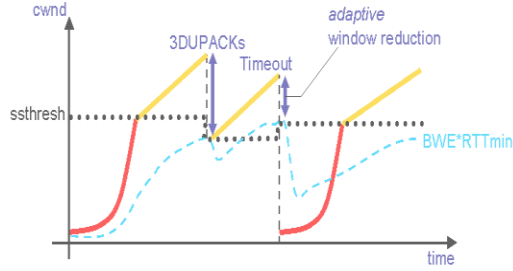


**30**  
Countries



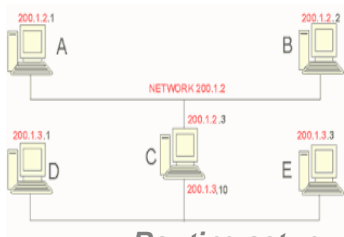
**100,000+**  
Experiments  
instantiated



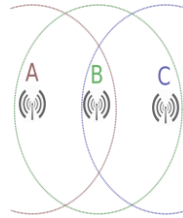


Congestion Avoidance (CA)  
Slow-start (SS)

*Protocol Behavior*

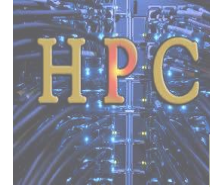


*Routing setup*



*Wireless Communication*

**Teach Basic Concepts**



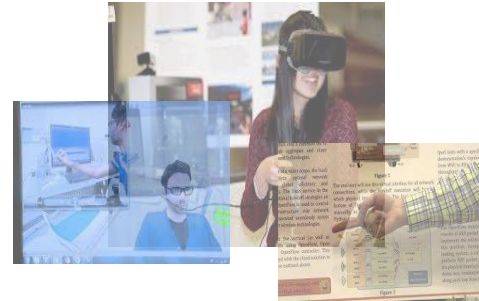
NFV

SDN

OpenFlow

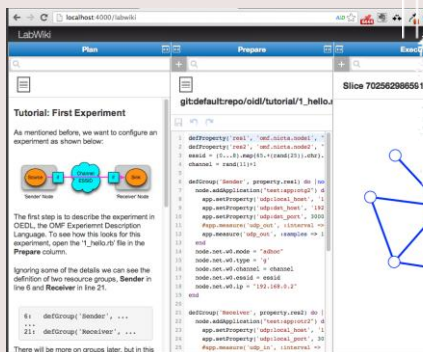


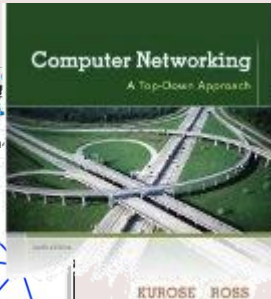
**New technologies**



**Semester Projects**



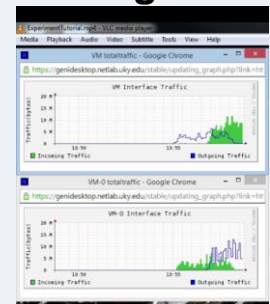




## Labs on GENI for networking textbook

Mike Zink  
UMass Amherst

## GENI Modules to teach networking concepts



*Example Demo Module*

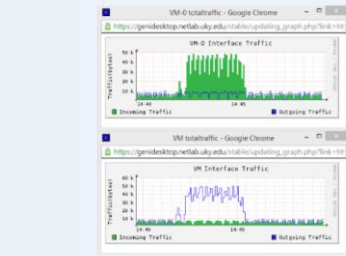


## Massive Online Open Courses on GENI

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



Shivendra Panwar, Thanasis Korakis  
NYU Poly



*Example Assignment*  
Kevin Jeffay, Jay Aikat  
UNC-Chapel Hill

# GENI Terminology



Sponsored by the National Science Foundation

**slice**

**project**

**experimenter**

**resource**

**aggregate**

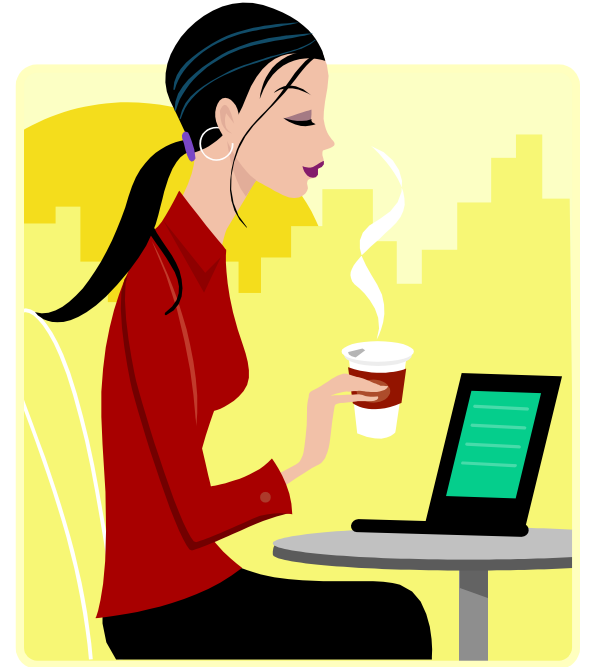


# An **experimenter**

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](http://www.geni.net)



# GENI User Authentication

The GENI Portal leverages InCommon  
for single sign-on authentication

**InCommon**<sup>®</sup>

Students from 518 educational  
and research institutions have  
InCommon accounts

For many experimenters:

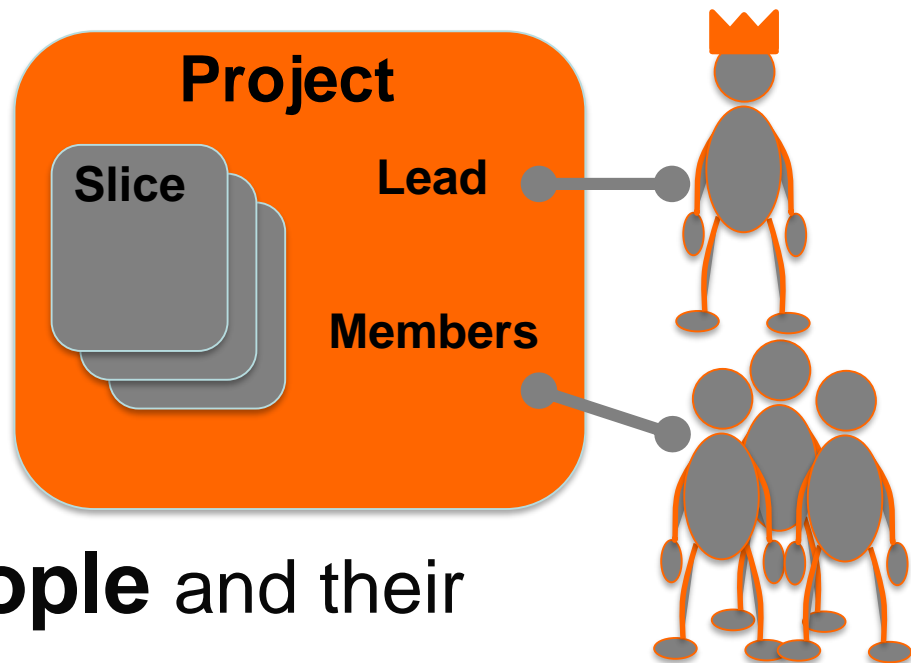
- no new passwords
- familiar login  
screens

**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**



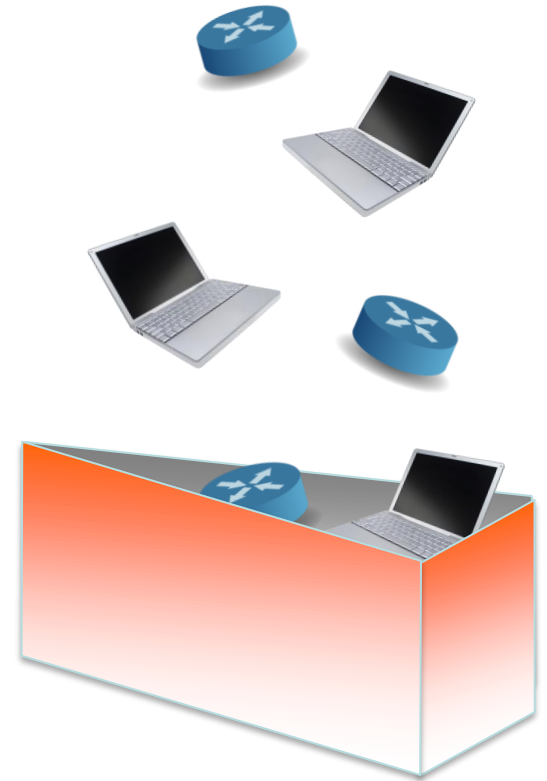
A **slice** is a *container* of resources used in an *experiment*.

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

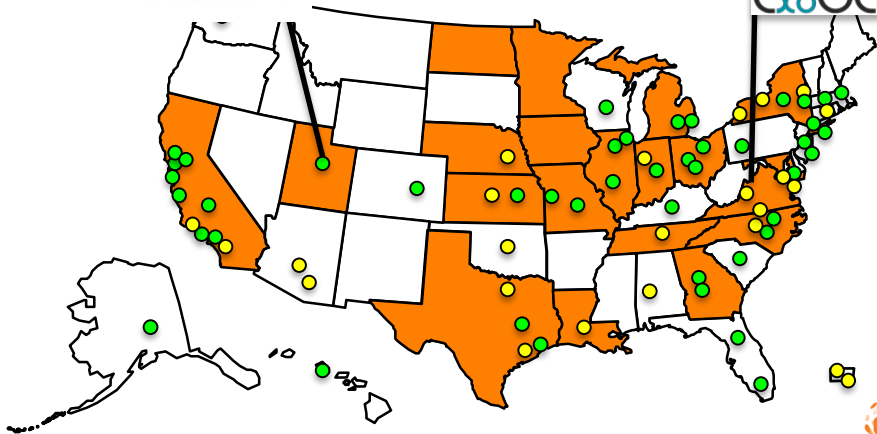
OpenFlow

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](http://www.geni.net)

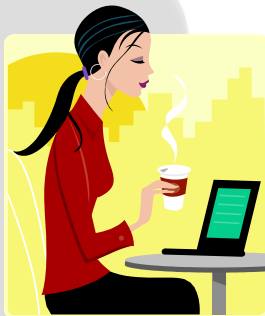




# SSH with a *password*

\*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
```



ssh



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

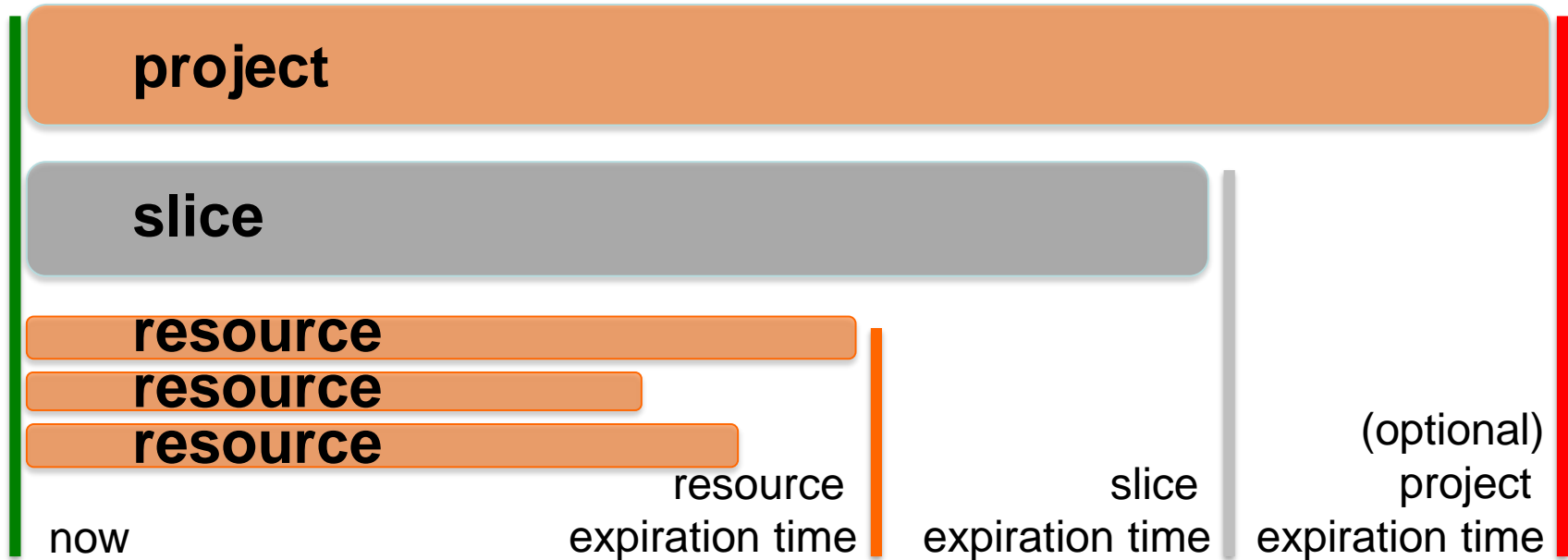


Sponsored by the National Science Foundation

# 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?

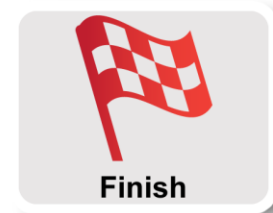
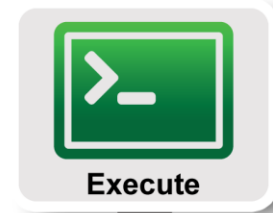
1. Grab a Worksheet and instructions
2. Did you do the pre-work?
  - A. Do you have an account?
  - B. Have you installed the tools?
    - \* SSH
3. Connect to the network

GENI Portal is at:

<http://portal.geni.net>

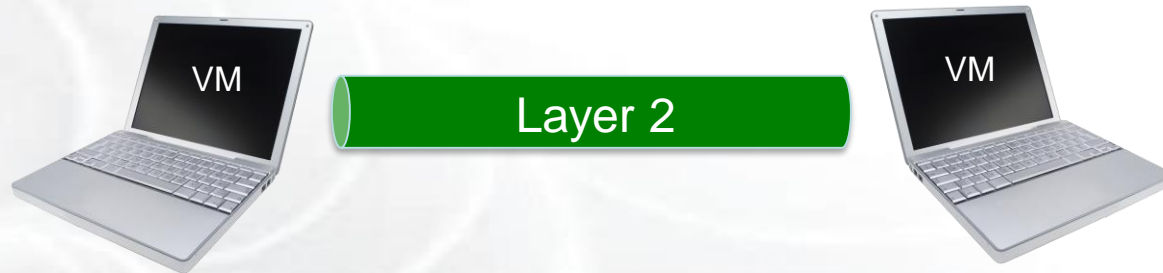
# Lab Zero: A First Experiment using GENI

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



## Do a Simple Experiment in GENI

Reserve two VMs connected at Layer 2





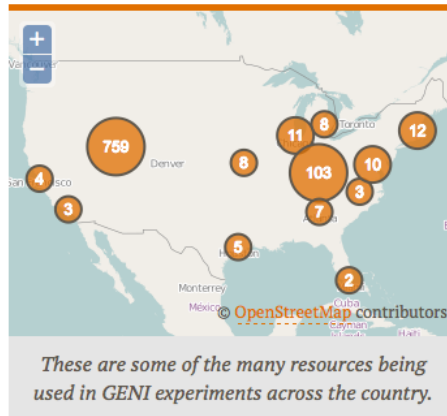
## WELCOME TO GENI

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](#)



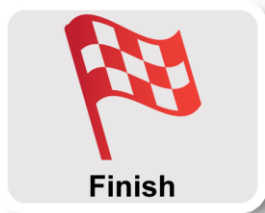
# Experiment Workflow



- Part I: Design/Setup



- Part II: Execute



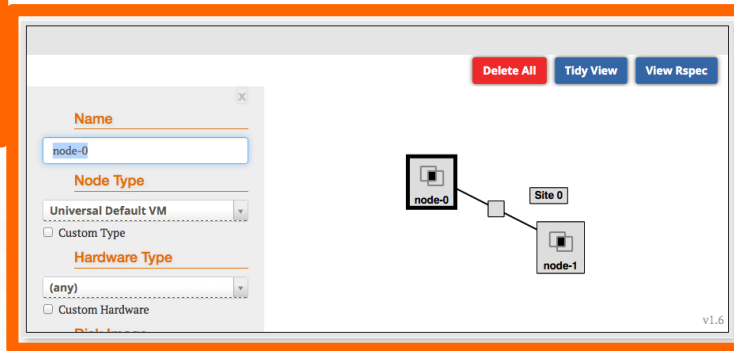
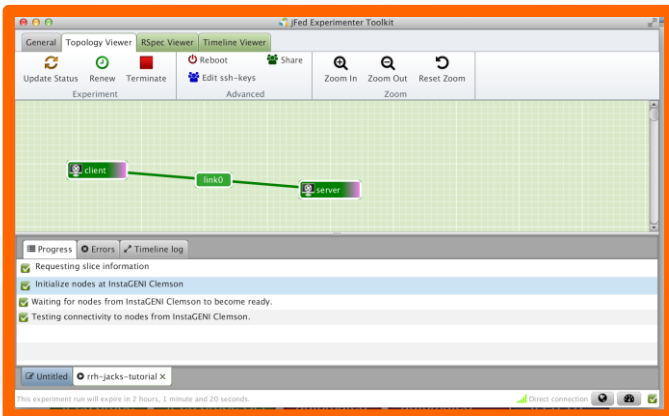
- Part III: Finish

# The GENI Portal is...

Use GENI

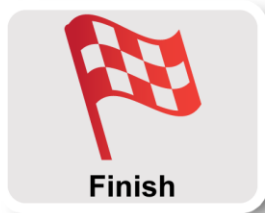
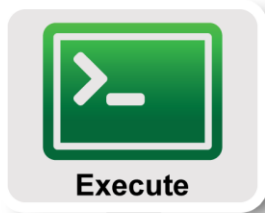
A web-based tool for experimenters to manage  
**experimenters, projects, and slices.**

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

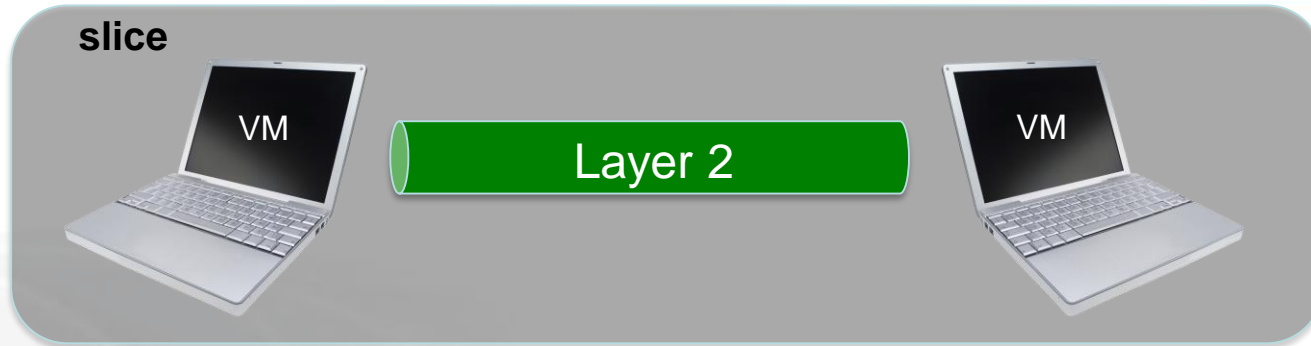
**Project Name: TrainTheTA**

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

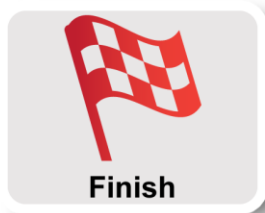
3.4 Check Whether VMs are Ready to be Used



- Part I: Design/Setup

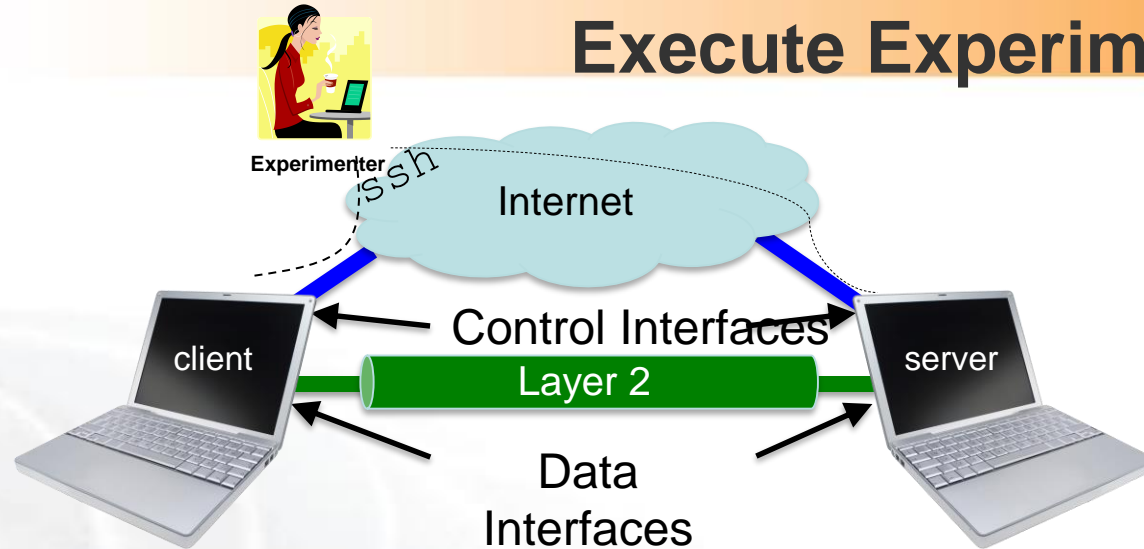


- **Part II: Execute**



- Part III: Finish

## Execute Experiment



4.1 Login to all three nodes

5.1 Test Connectivity

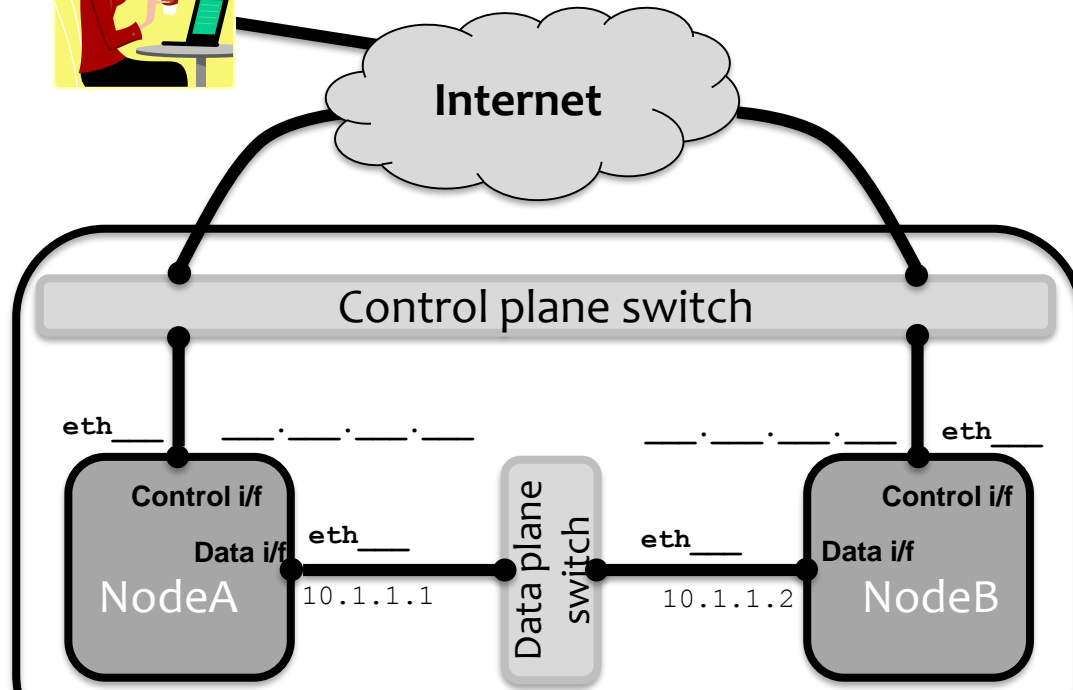
5.2 Explore the Data and Control Planes

6.1 Logout of nodes

# Worksheet

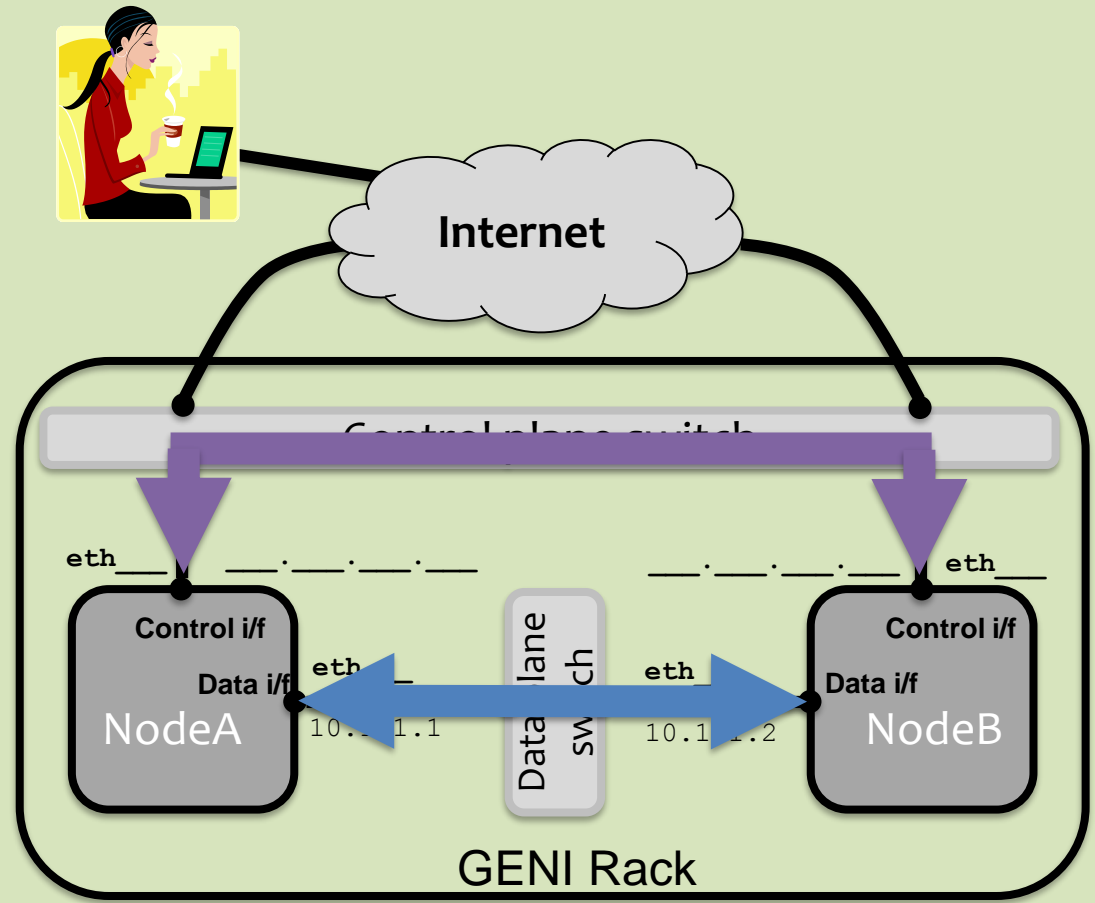
Slice Name:

lab0<your initials>

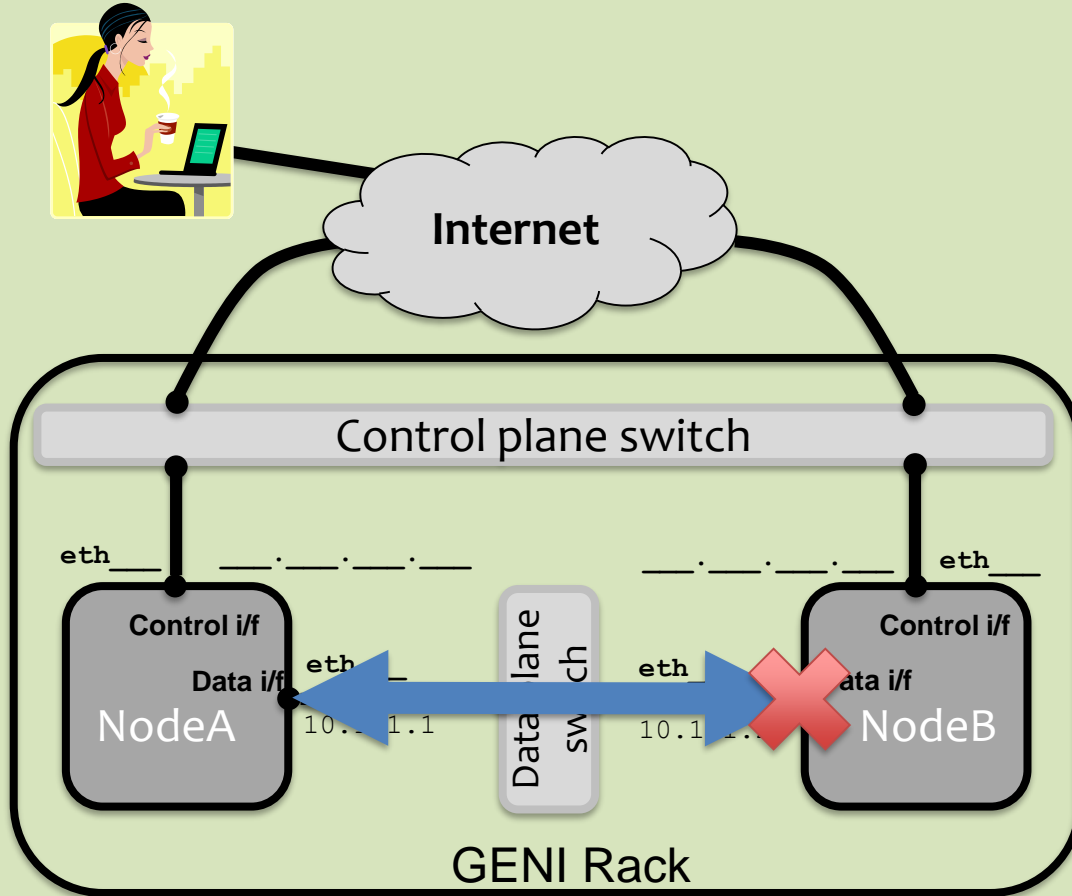


5.1

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

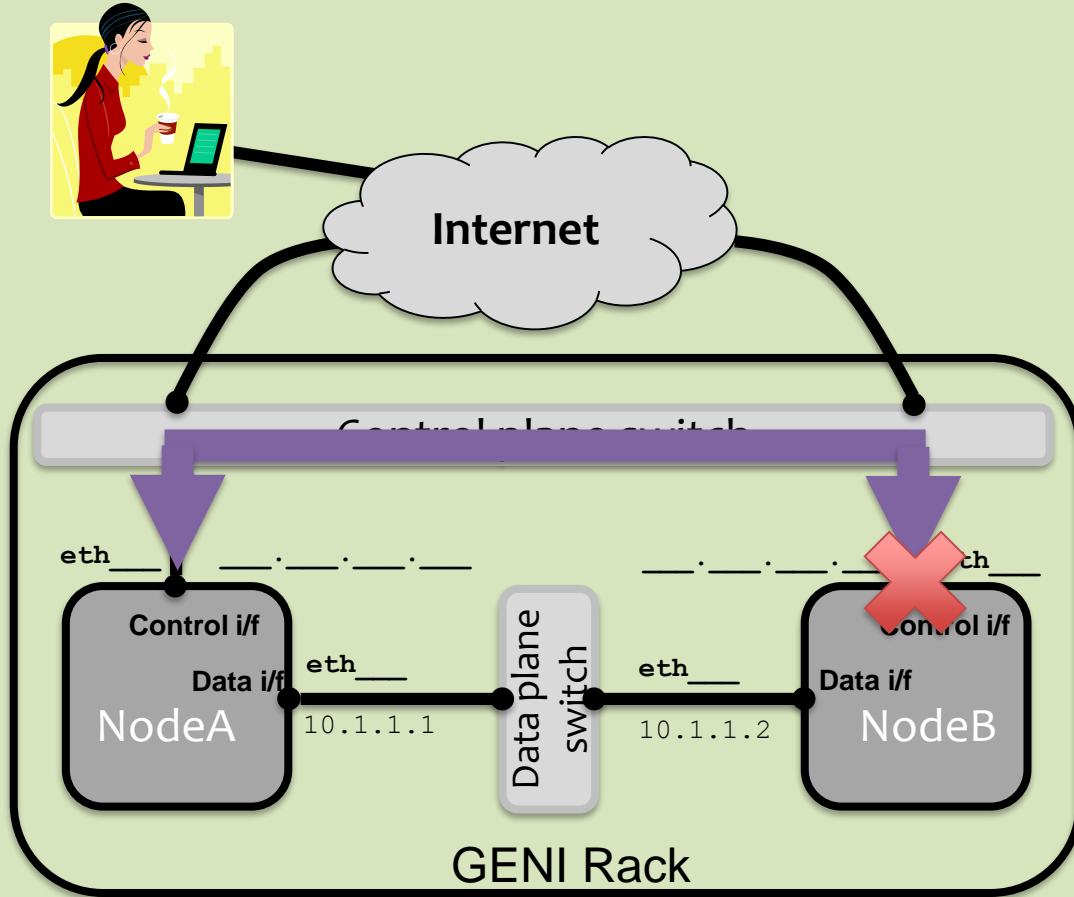


5.3 When you bring down the **data** interface, the destination should become unreachable. Why?

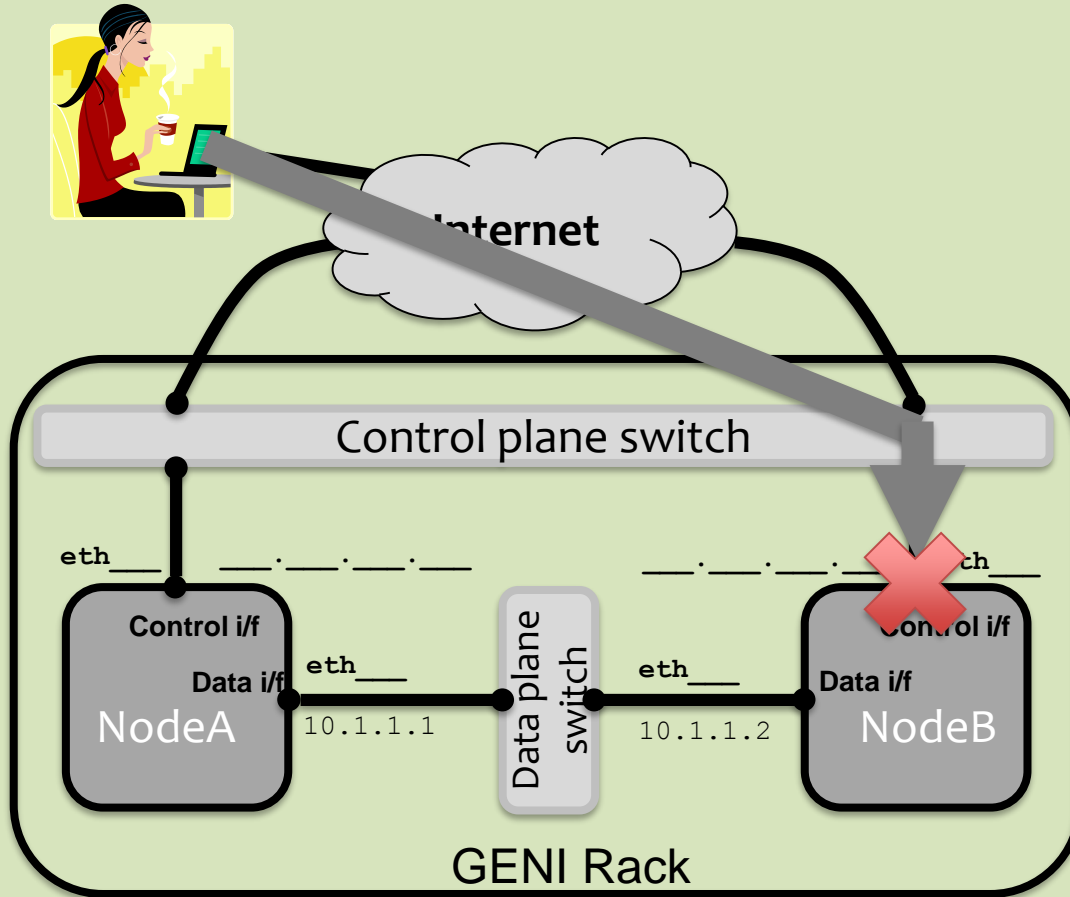


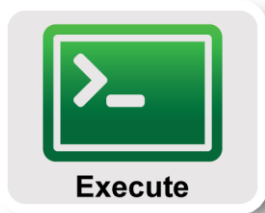


5.3 After you bring down the **control** interface, the destination becomes unreachable. Why?



- 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**

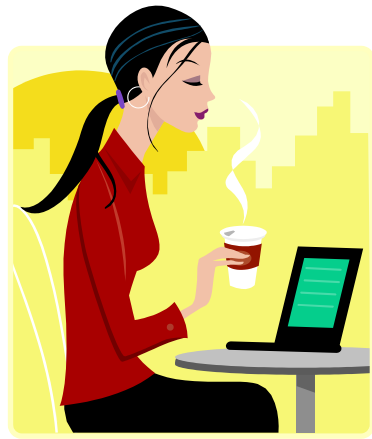
GENI Resource Specifications

**GENI AM API**

GENI Aggregate Manager API



Sponsored by the National Science Foundation



researcher tool



create slice

*CH APIs*

clearinghouse



slice credential

*slice, RSpec*

*Aggregate Manager API*



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

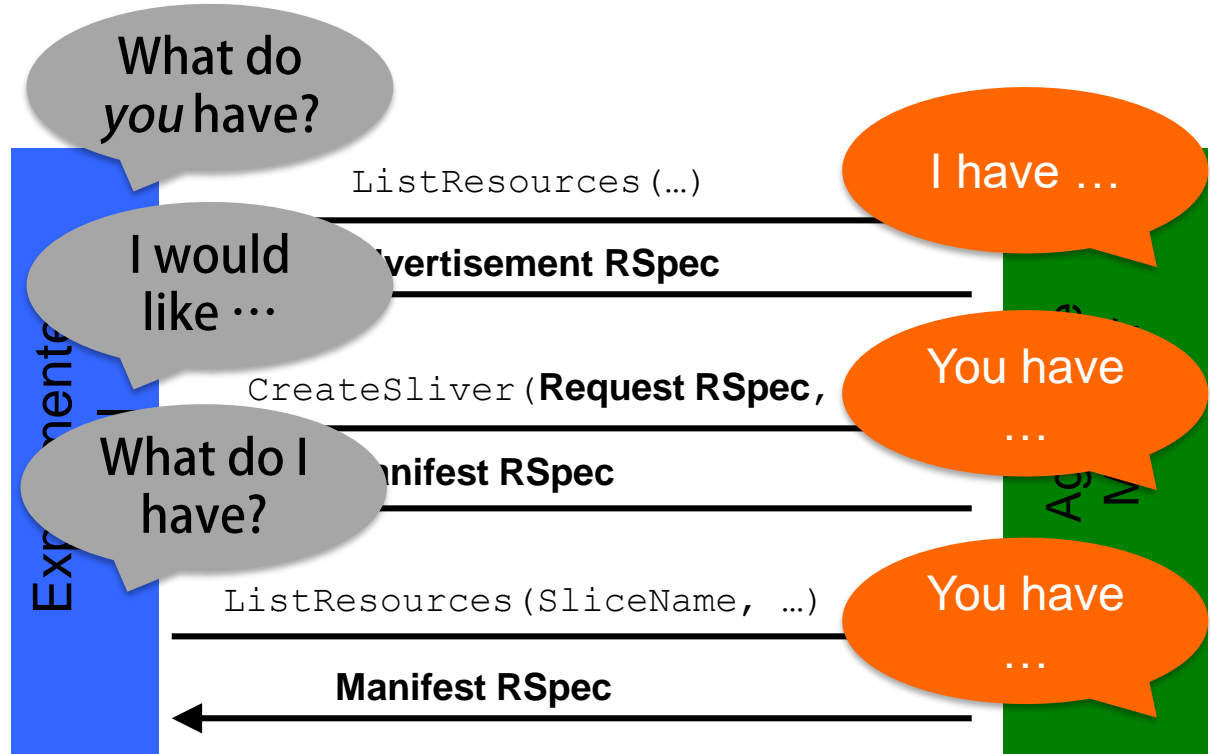
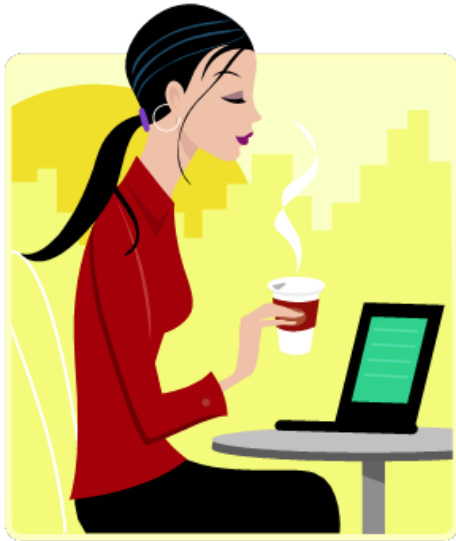
## Manifest RSpecs

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

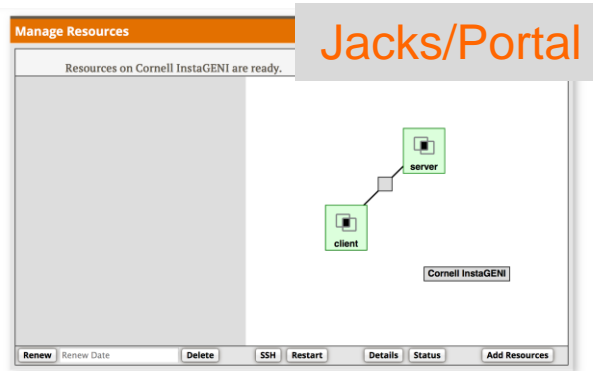
[www.geni.net](http://www.geni.net)





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

Hosted



The screenshot shows the 'Manage Resources' page of the Jacks/Portal. It features a header with the title 'Manage Resources' and a sub-header 'Resources on Cornell InstaGENI are ready.' Below this is a diagram showing a 'client' node connected to a 'server' node, with a 'Cornell InstaGENI' label below the connection. At the bottom, there is a toolbar with buttons for 'Renew', 'Renew Date', 'Delete', 'SSH', 'Restart', 'Details', 'Status', and 'Add Resources'.

Jacks/Portal

Local

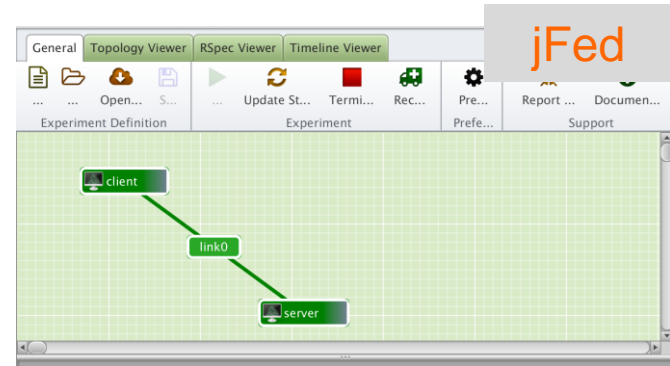
```
$ omni createsliver aliceslice myRS
INFO:omni: -----
INFO:omni: Completed createsliver:

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. It has a menu bar with options: 'General', 'Topology Viewer', 'RSpec Viewer', and 'Timeline Viewer'. Below the menu is a toolbar with icons for 'Open...', 'Update St...', 'Termi...', 'Rec...', 'Pre...', 'Report ...', and 'Documen...'. The main area displays a network diagram with a 'client' node connected to a 'server' node via a 'link0' link. The background is a light green grid.

jFed



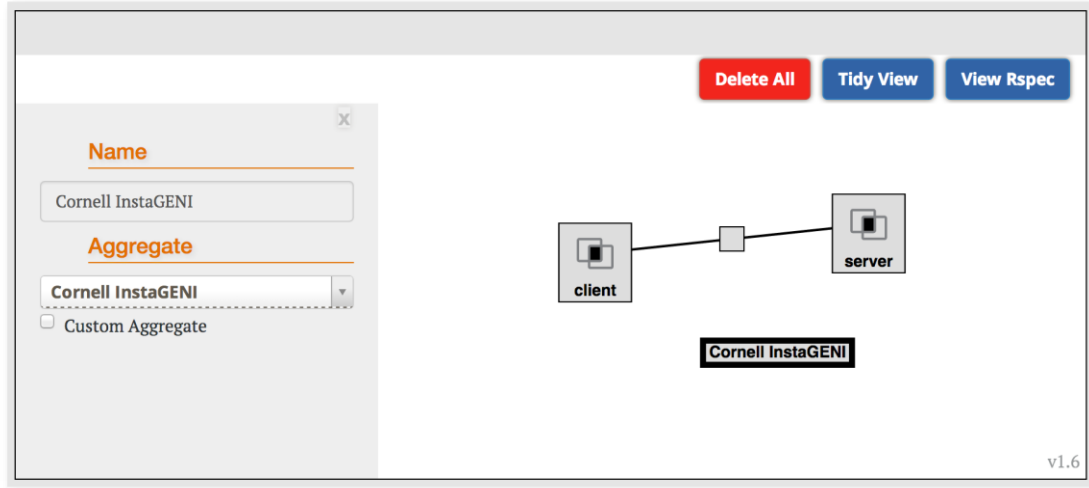
# GENI Tool: Jacks in the Portal

## *A graphical user interface*



Sponsored by the National Science Foundation

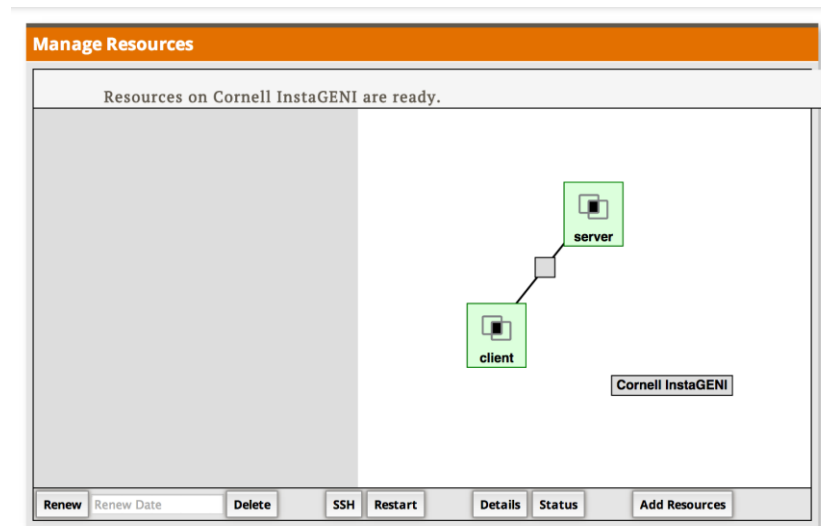
When you draw a topology using a tool, it creates a **request RSpec** describing the resources

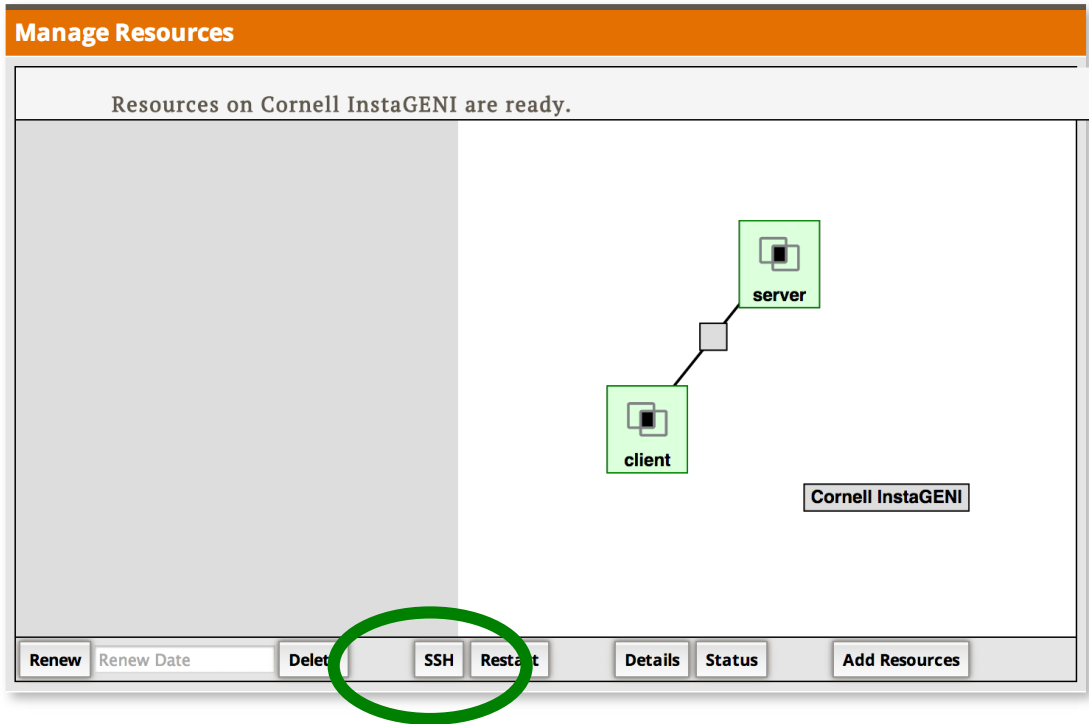


When you reserve resources at an aggregate, **createSliver** is called using that request

Tools periodically call `sliverStatus` at each used aggregate

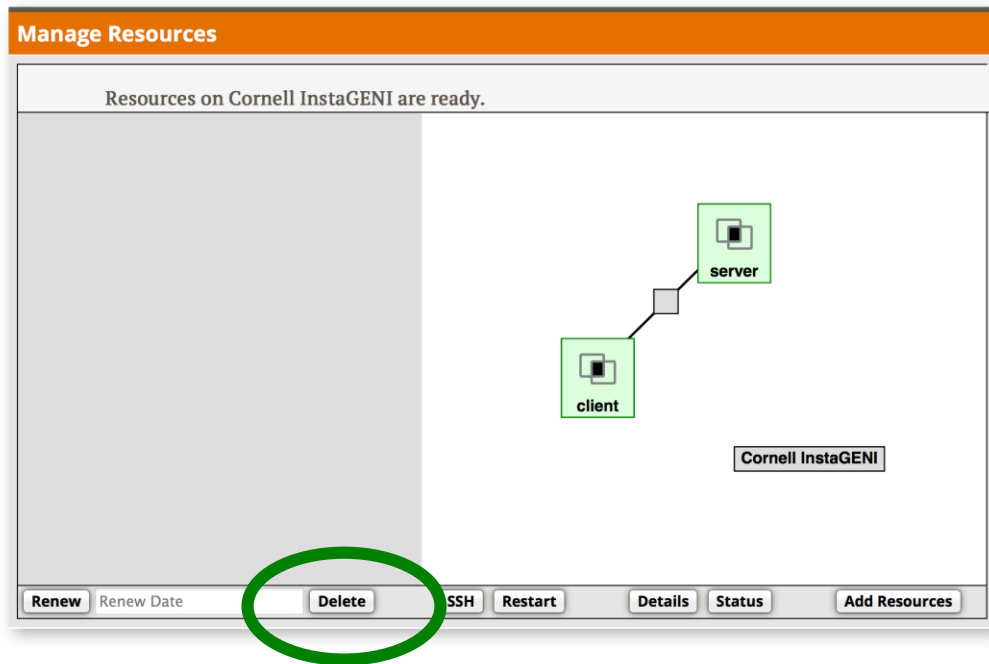
When the returned status is “ready”, the resources turn green





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*



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# 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

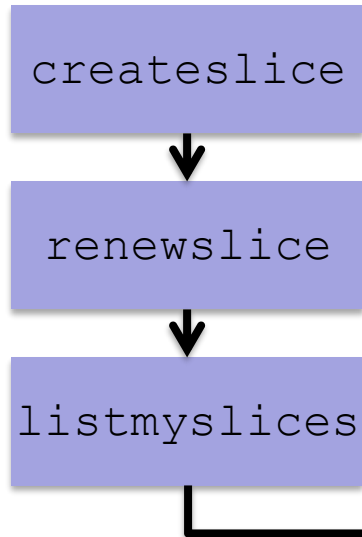
```
Result Summary: Slice urn:publicid:IDN+pgeni
Reserved resources on https://www.emulab.net/p
Saved createsliver results to aliceslice-man
INFO:omni: =====
```





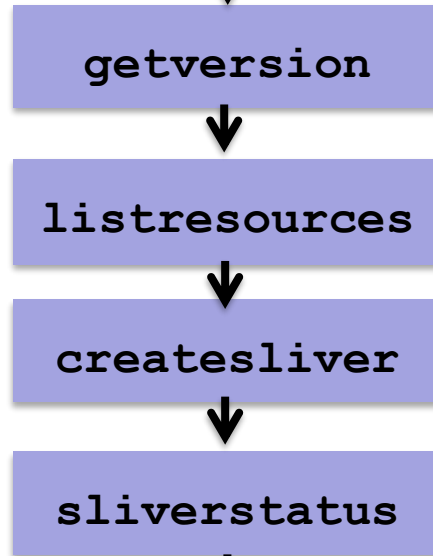
# omni Workflow

## 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`



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# GENI Support for Automating Resource Reservation



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# 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

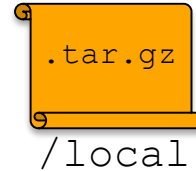
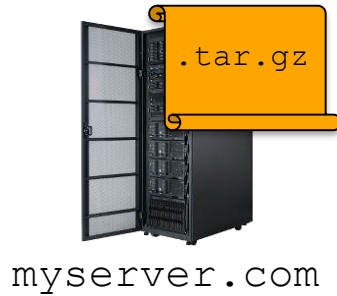


# Install and Execute Scripts





# 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*



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[www.geni.net](http://www.geni.net)



## Next Session

### 3:00 PM ET on Fri, Feb 10

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:  
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