

TEST-DRIVEN EVALUATION OF GALAXY SCALABILITY ON THE CLOUD

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Overview

- GVL Quality Assurance
 - Automated QA tests

- GVL Scaling
 - Many QA tests in parallel = Scalability test



Quality Assurance

Why?

With each new GVL release:

- Do the tutorials run to completion? (Tedious and error prone to check manually)
- Need a quick way of knowing whether things are in reasonable shape
- Need an end-user perspective on how things work

How?

- Using Selenium
- Run full workflows that exercise a complete set of tools
- Check whether tool output == expected output
- Also exercise typical use cases in UI



Demo

Selenium in action

Se Grid Console v.2.40.0	, Help
DefaultRemoteProxy (version : 2.40.0) id : http://115.146.86.191:5555, OS : LINUX	DefaultRemoteProxy (version : 2.40.0) id : http://115.146.86.82:5555, OS : LINUX
Browsers Configuration	Browsers Configuration
Remote Control (legacy) v: v: v: v: v: v: v: v:	Remote Control (legacy) v: v: v: v: v: v: v: v:
DefaultRemoteProxy (version : 2.40.0) id : http://115.146.86.192:5555, OS : LINUX	DefaultRemoteProxy (version : 2.40.0) Id : http://115.146.86.79:5555, OS : LINUX
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DefaultRemoteProxy (version : 2.40.0) id : http://115.146.85.159:5555, OS : LINUX Browsers Configuration	DefaultRemoteProxy (version : 2.40.0) Id : http://115.146.86.76:5555, OS : LINUX Browsers Configuration
Remote Control (legacy) v: € v: € ● v: ● ● • ●	Remote Control (legacy) v: v: Image: Second Seco

view config





Writing a simple test

```
1⊖ from gvl_test_base import GVLTestBase
2 from selenium_snippets.galaxy.history import History
3 from selenium_snippets.galaxy.get_data import GetData
   from selenium_snippets.galaxy.rna_analysis import RNAAnalysis
4
   from selenium_snippets.galaxy import snippet_base
5
6
7⊖ class SimpleTest(GVLTestBase):
8
90
       def __init__(self, galaxy_test_context):
           super(SimpleTest, self).__init__(galaxy_test_context)
10
11
12
       @snippet_base.ui_action()
13⊝
       def execute_gvl_testcase(self):
14
           history = History(self.context)
15
           history.create_new_history("Hello world")
16
           file = "https://swift.rc.nectar.org.au:8888/v1/AUTH_377/public/RNAseqDGE_BASIC/C2_R1.chr4.fq"
17
           GetData(self.context).run_upload_file('url', file, 'fastgsanger')
18
           history.wait_for_datasets_to_finish()
19
           RNAAnalysis(self.context).run_tophat(file, "D melanogaster (dm3)")
20
           history.wait_for_datasets_to_finish()
21
```

A library of test snippets available. Composed as desired for a more complex test. Similar to Galaxy's Twill based tests of internal API



Issues encountered

Some elements have no ids = brittle xpaths



Performance Testing

Why?

- How many workers do you need for an RNASeq wokshop with 20 users? What size should the workers be?
- How does the GVL scale for different workloads?
- What combinations of storage, instance types, workers etc. are recommended?
- We had mostly anecdotal evidence needed a more data-driven approach

How?

- 1 thread = 1 user = QA
- Many threads = Multiple Users = Performance
- Use Selenium Grid



Selenium Grid



Also tried PhantomJS+Ghostdriver as a lightweight Selenium backend – lots of potential – but didn't work out



What was done?

- Desirable combinations tested.
- All run on the NCI zone (identical hardware)
- Each test had independent resources (e.g. brand new Galaxy/Cloudman instance launched, independent gluster servers, nfs servers used etc.)
- Transient cloud conditions not controlled for



Combinations tested

- storage_type = { gluster, transient, volumes, nfs }
- machine_type = { m1.medium, m1.large, m1.xlarge }
- workers = { 0 to 5 workers }
- workloads = { rnaseq basic tutorial, deseq basic tutorial microbial assembly tutorial, variant detection basic tutorial }
- simultaneous users = { 1, 5, 10, 20 }

Test loop

for storage_type in storage_types: (4) for machine_type in machine_types: (3) for worker in workers: (5) for workload in workloads: (4) for user in number_of_users: (4) time_stuff()

- Total = 4 * 3 * 6 * 4 * 4 = 1152
- Total completed so far: 837
- Successful completion for: 655
- Reasons for failure:
 - Turnaround time for a job capped at 1.5 hours
 - Transient capacity issues on the cloud (couldn't get the machines on demand)
 - The occasional selenium hiccup



What it records

• Time taken for each segment of the test



5 GB of atop logs and timing logs (mostly atop)

- Records atop logs at 10 second intervals
 - Provides snapshot of CPU, memory, process and network usage

Results

A list of configurations to use for a particular scenario (e.g. how many workers for a 20 user rna-seq workshop?)

Average of time taken workers	machine_type 1. m1.medium	2. m1.large	3. m1.xlarge
0	1425.393948	1569.968806	1266.41437
1	1346.412998	1176.365253	1084.031046
2	1269.079958	1124.207206	993.8295797
3	1938.757942	1380.971698	937.1919329
4	1558.144051	1133.047089	819.9848138
5	1177.656928	986.7495938	722.8780805





 One worker pays off the most irrespective of the instance type or workload





- Transient vs Volumes
 - Volumes were slightly outperforming transient storage.
 - We asked the NeCTAR team why? Turned out that transient storage was rate limited to 25MB/sec to prevent any one VM from hogging the disk bandwidth.





- Gluster vs Volumes
 - Differences turned out to be marginal.
 - Not congruent with our previous experiences
 - Reasons unknown so far



- Total CPUs in cluster appears to contribute most to overall performance.
 - E.g. Two large (4 core) instances roughly = Single xlarge (8 core) instance
- Therefore less likely to overprovision if you use many smaller instances (with autoscaling), as opposed to a few larger instances





Repository

https://bitbucket.org/gvl/gvl-stress-test

Raw Result Data:

url: <u>https://swift.rc.nectar.org.au:8888/v1/AUTH_377/gvl_performance_results</u> shortened url: <u>http://bit.ly/gvl_performance_results</u>

Detailed Report:

Work in progress



What next?

- Amazon/EC2 vs NeCTAR/Openstack?
- Gluster vs NFS vs PVFS/OrangeFS vs …?
- No. of web runners?
- No. of Job Handlers?
- No. of Nginx workers?
- More in-depth analysis of the data we have right now.