



How to set up a Galaxy service you can count upon

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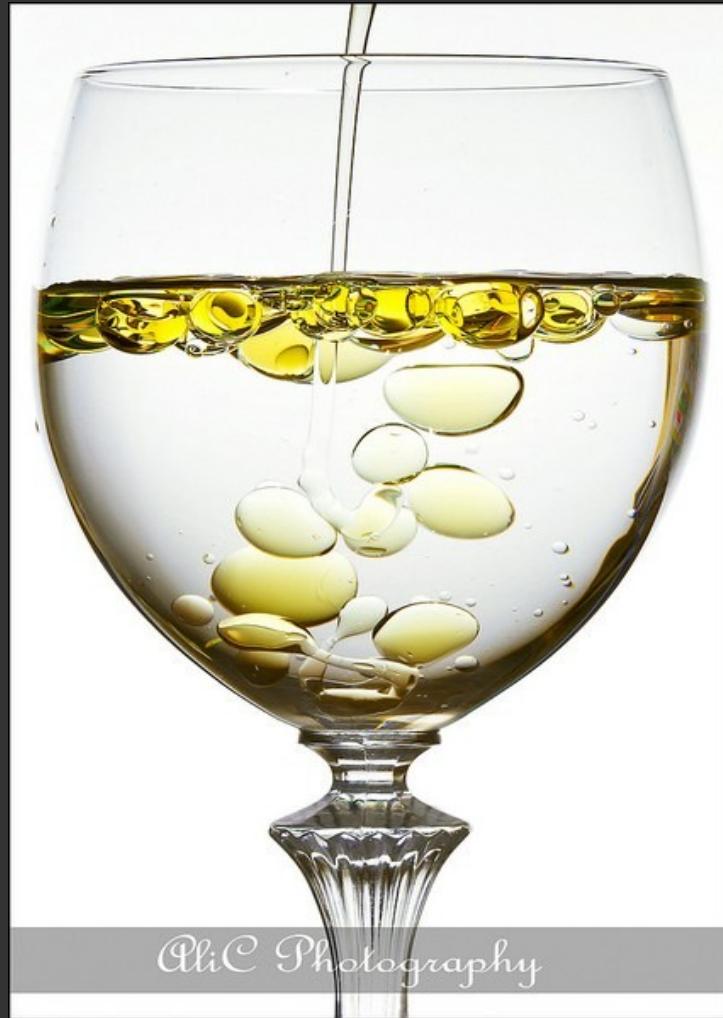


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The Zen of Galaxy

- ➊ Keep your modifications isolated
- ➋ Use virtualenv
- ➌ Test
- ➍ Test more !
- ➎ Use scalable storage/cluster
- ➏ Know what Galaxy is doing



AliC Photography

Don't mix your code & configuration with Galaxy

HOW TO :

- Your xml configuration goes here :

universe_wsgi.ini :

tool_config_file=tool_conf.xml,shed_tool_conf.xml,**my_own_confs.xml**

- Your app's folder structure :

tools/

myApp/

 myapp.py

 myapp.xml

- MyApp → my_own_conf.xml

Use *virtualenv*

- ➊ Virtualenv : for isolated *Python* environments
- ➋ Control your libraries
- ➌ Each virtualenv has its own pip/easy_install
- ➍ pip freeze > requirements.txt

Test !

- `run_unit_tests.sh`
- `run_functional_tests.sh`
- `run_tool_shed.sh`
- `run_tool_shed_functional_tests.sh`

- Use scalable storage/cluster

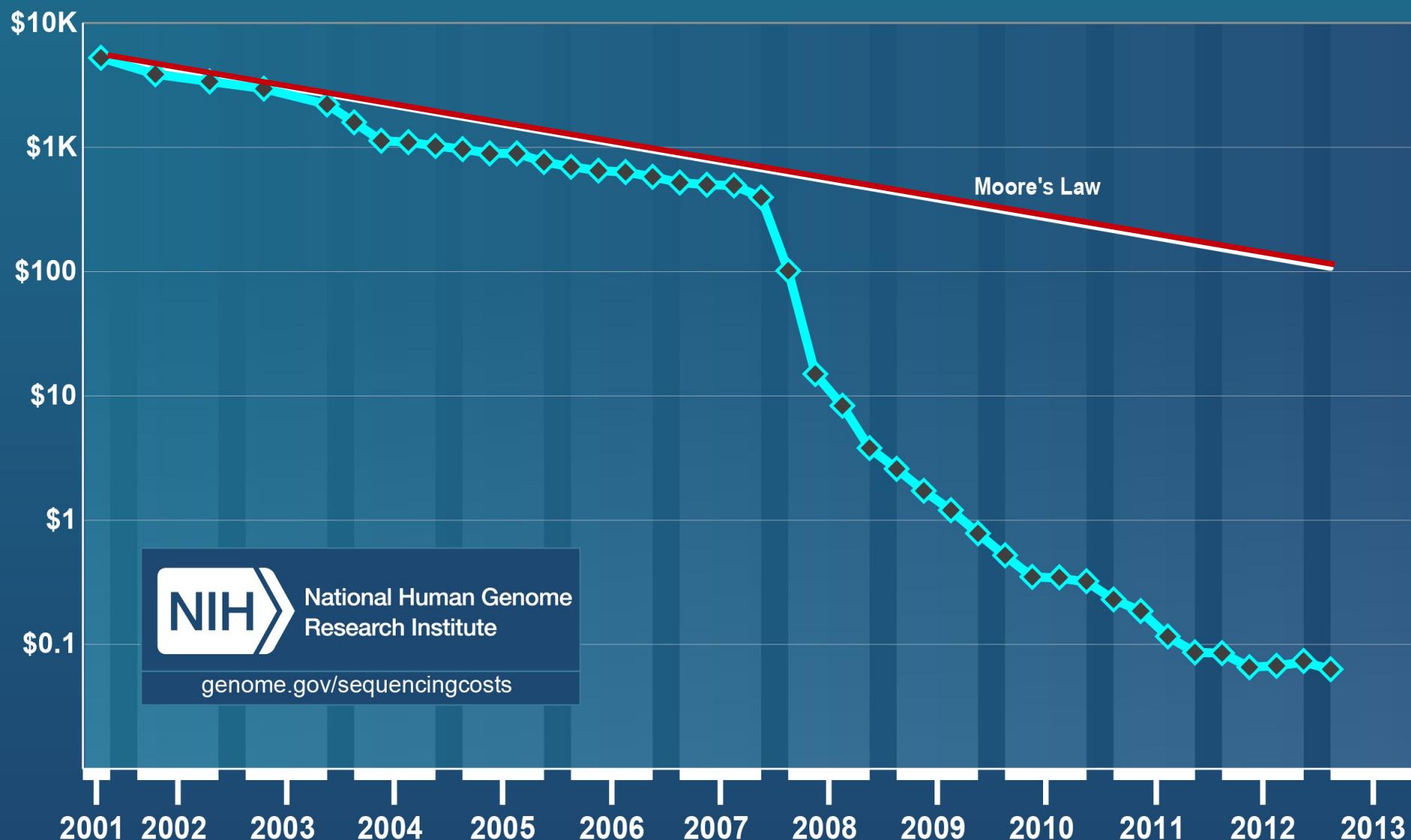
- ◎ ZFS rocks !

- ◎ some theoretical limits in ZFS are :

- ◎ 2^{48} : number of entries in any individual directory
- ◎ 16 Exbibytes (2^{64} bytes): maximum size of a single file
- ◎ 16 Exbibytes: maximum size of any attribute
- ◎ 256 Zebibytes (2^{78} bytes): maximum size of any zpool
- ◎ 2^{56} : number of attributes of a file (actually constrained to 2^{48} for the number of files in a ZFS file system)
- ◎ 2^{64} : number of devices in any zpool
- ◎ 2^{64} : number of zpools in a system
- ◎ 2^{64} : number of file systems in a zpool

- ◎ Snapshots, deduplication, compression & more

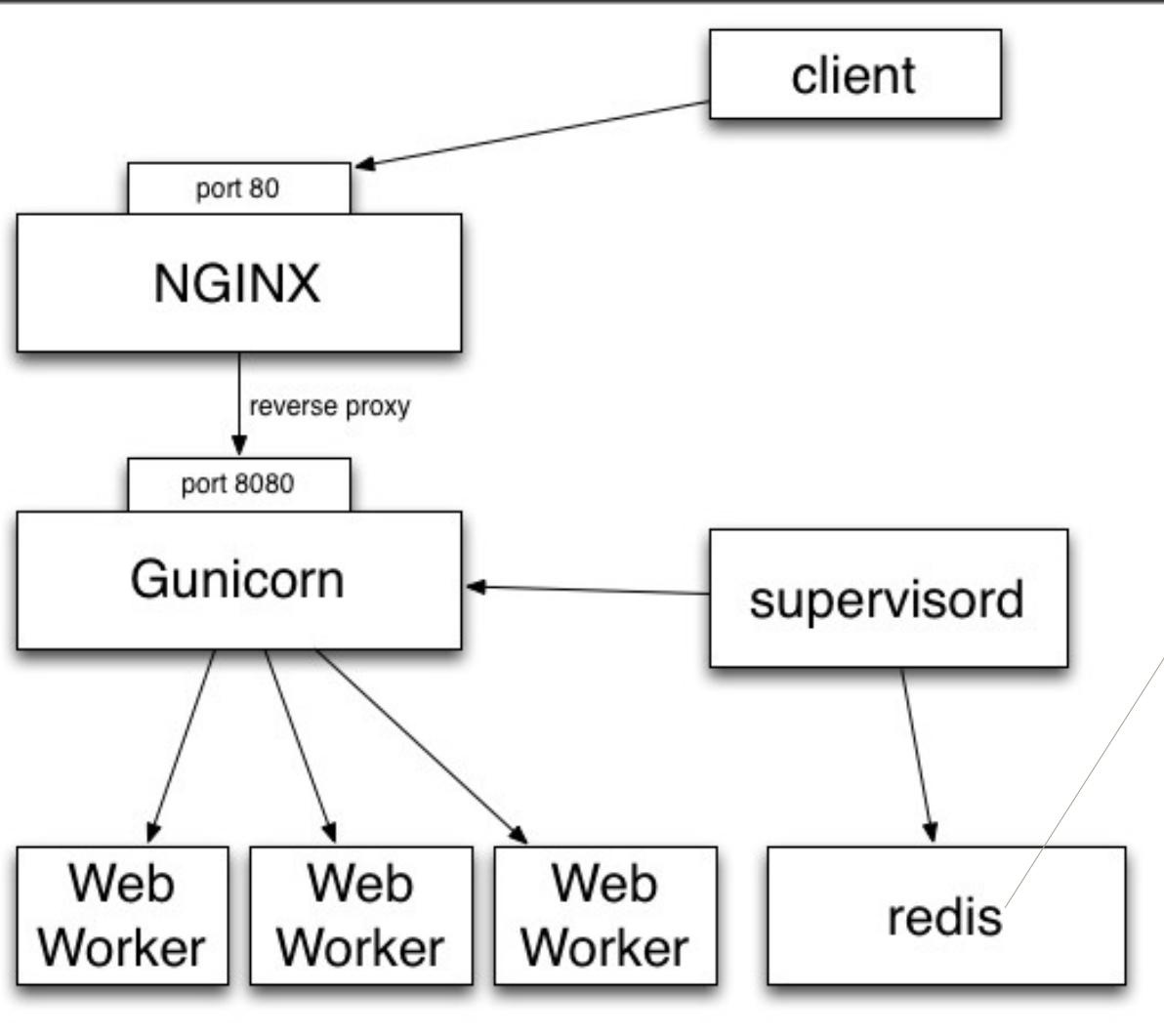
Cost per Raw Megabase of DNA Sequence



National Human Genome
Research Institute

genome.gov/sequencingcosts

Web stack the python way (The GIL issue)



Dans Galaxy : universe_wsgi.ini

⌚ Sections :

⌚ server (web)

⌚ Manager

⌚ Handler

```
[server:web0]
use = egg:Paste#http
port = 8080
host = 127.0.0.1
use_threadpool = true
threadpool_workers = 7
```

```
[server:web1]
use = egg:Paste#http
port = 8081
host = 127.0.0.1
use_threadpool = true
threadpool_workers = 7
```

```
[server:handler0]
use = egg:Paste#http
port = 8090
host = 127.0.0.1
use_threadpool = true
threadpool_workers = 5
```

```
[server:handler1]
use = egg:Paste#http
port = 8091
host = 127.0.0.1
use_threadpool = true
threadpool_workers = 5
```

Monitor your Galaxy

⌚ What is Circus ?

⌚ process manager

- ⌚ Starts & stops processes

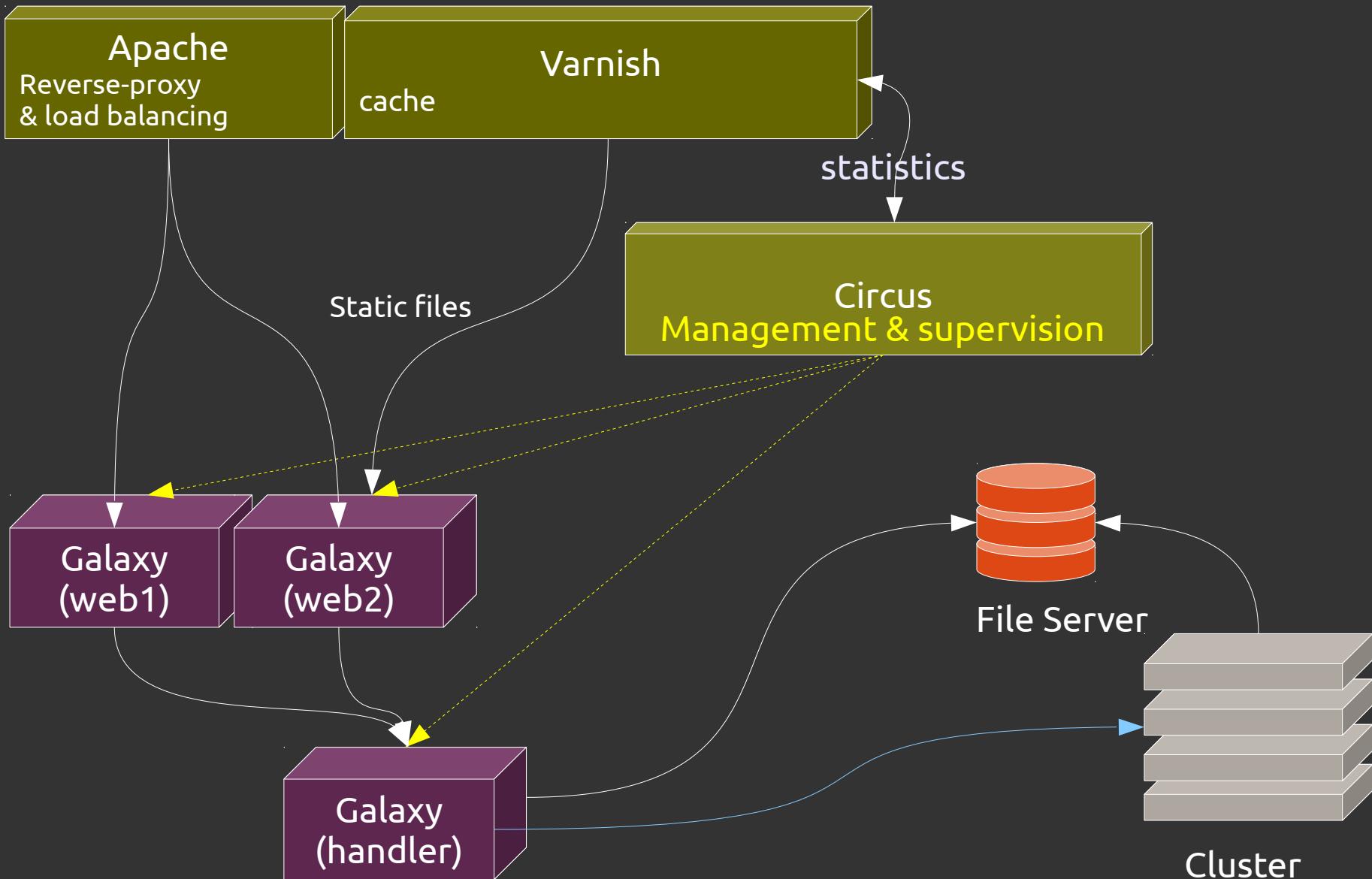
- ⌚ Automatic « respawn », contrôle du « flapping »

⌚ monitoring

⌚ Real time statistics

- ⌚ ZeroMQ → asynchronous communication

Notre instance Galaxy

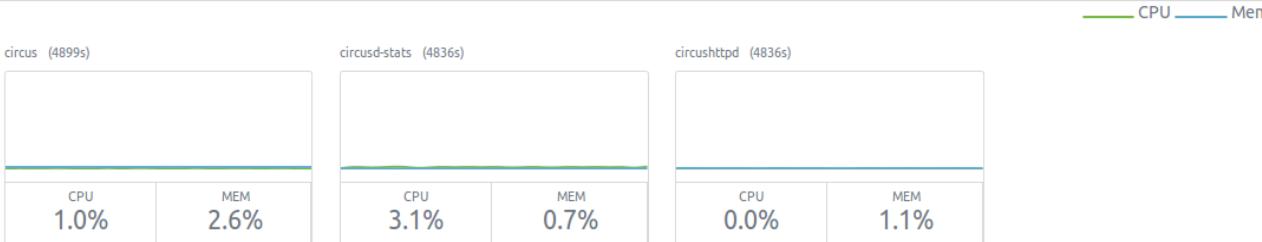


[Add Watcher](#)

Watchers

Name Status	Processes	Command	Shell	uid	gid
handler0	1	python ./scripts/paster.py serve universe_wsgi.ini -server-name=handler0 -pid-file=/srv/galaxy/galaxy-dist/run/galaxy-handler0.pid -log-file=/srv/galaxy/galaxy-dist/log/galaxy-handler0.log	False	None	None
main	1	python ./scripts/paster.py serve universe_wsgi.ini -server-name=main -pid-file=/srv/galaxy/galaxy-dist/run/galaxy-main.pid -log-file=/srv/galaxy/galaxy-dist/log/galaxy-main.log	False	None	None
manager	1	python ./scripts/paster.py serve universe_wsgi.ini -server-name=manager -pid-file=/srv/galaxy/galaxy-dist/run/galaxy-manager.pid -log-file=/srv/galaxy/galaxy-dist/log/galaxy-manager.log	False	None	None
web1	1	python ./scripts/paster.py serve universe_wsgi.ini -server-name=web1 -pid-file=/srv/galaxy/galaxy-dist/run/galaxy-web1.pid -log-file=/srv/galaxy/galaxy-dist/log/galaxy-web1.log	False	None	None

Circus Daemons



Options

ZeroMQ endpoint used to connect clients like <i>circusctl</i> or <i>circushttpd</i>	<code>tcp://127.0.0.1:5555</code>
ZeroMQ endpoint used for pub/sub	<code>tcp://127.0.0.1:5556</code>
ZeroMQ endpoint used for the pub/sub stats	<code>tcp://127.0.0.1:5557</code>
Delay in seconds for the ZMQ polling.	1.0

