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# My SolrCloud Experience



- Currently, working on scaling up to a 200+ node deployment at LucidWorks
- Operated 36 node cluster in AWS for Dachis Group (1.5 years ago, 18 shards ~900M docs)
- Contributed several tests and patches to the code base
- Built a Fabric/boto framework for deploying and managing a cluster in EC2
- Co-author of Solr In Action; wrote CH 13 which covers SolrCloud

# Solr Scaling Toolkit



- Requirements
- High-level overview
- Nuts and Bolts (live demo)
- Roadmap
- Q&A

#### Tasks to Automate



- Provisioning N machine instances in EC2
- Configuring / starting ZooKeeper (1 to n servers)
- Configuring / starting N Solr instances in cloud mode (M x N nodes)
- Integrating with Logstash4Solr and other supporting services, e.g. collectd
- Day-to-day operations on an existing cluster

## Python-based Tools



**boto** - Python API for AWS (EC2, S3, etc)

**Fabric** - Python-based tool for automating system admin tasks over SSH

**pysolr** - Python library for Solr (sending commits, queries, ...)

kazoo - Python client tools for ZooKeeper

#### **Supporting Cast:**

JMeter - run tests, generate reports collectd - system monitoring Logstash4Solr - log aggregation JConsole/VisualVM - monitor JVM during indexing / queries

#### Fabric in 3 minutes or Less ...



Fabric helps you do common system administration tasks on multiple hosts over SSH ...

- Just Python
- Easy to install / learn; good documentation
- http://docs.fabfile.org/en/1.8/

### Fabric in 3 minutes or Less, cont. ...



- Define all commands in a file named: fabfile.py
- Get a list of supported commands with short description

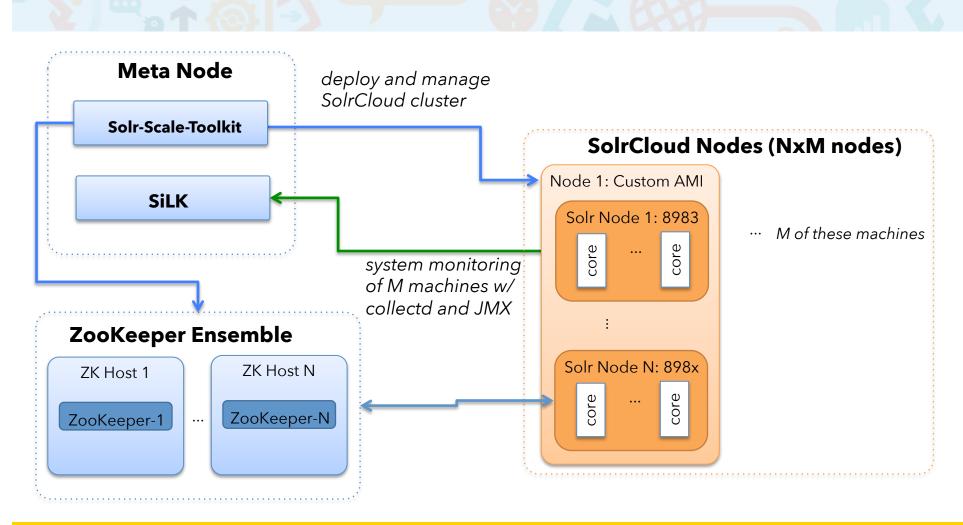
```
$ fab -1
Available commands:
backup_to_s3
check_zk
commit
Sends a hard commit to the ...
```

Get extended documentation for a command

```
$ fab -d new_solr_cloud
Displaying detailed information for task 'new_solrcloud':
    Provisions n EC2 instances and then deploys SolrCloud; uses
    the new_ec2_instances and setup_solrcloud commands ...
```

#### Solr Scale Toolkit: Architecture





## Provisioning machines



#### fab new\_ec2\_instances:test1,n=3,instance\_type=m3.xlarge

- Custom built AMI?
- Block device mapping
  - dedicated disk per Solr node
- Launch and then poll status until they are live
  - verify SSH connectivity
- Tag each instance with a cluster ID and username

### ZooKeeper

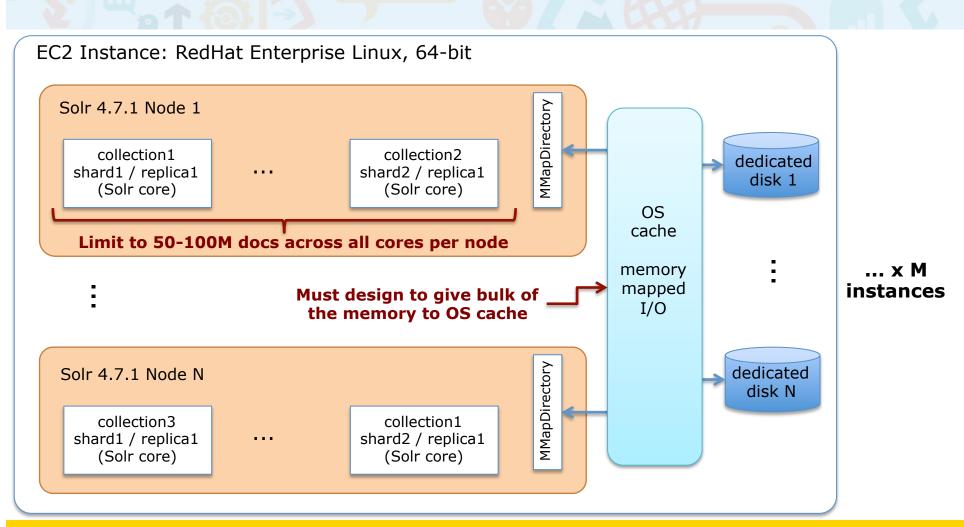


# fab new\_zk\_ensemble:zk1,n=3

- Two options:
  - provision 1 to N nodes when you launch Solr cluster
  - use existing named ensemble
- Fabric command simply creates the myid files and zoo.cfg file for the ensemble
  - and some cron scripts for managing snapshots
- Basic health checking of ZooKeeper status:
  - echo srvr | nc localhost 2181

### SolrCloud Cluster: NxM nodes





#### SolrCloud



## fab new\_solrcloud:test1,zk=zk1,nodesPerHost=2

- Upload a BASH script that starts/stops Solr
- Set system props: jetty.port, host, zkHost, JVM opts
- One or more Solr nodes per machine
- JVM mem opts dependent on instance type and # of Solr nodes per instance
- Optionally configure log4j.properties to append messages to Rabbitmq for Logstash4Solr integration

### solr-ctl.sh



- BASH script that implements:
  - start/stop Solr nodes on each EC2 instance
  - sets JVM memory options, system properties (jetty.port), enable remote JMX, etc
  - backup log files before restarting nodes
  - ensure JVM is killed correctly before restarting
- Environment variables in:
  - solr-ctl-env.sh

## Miscellaneous Utility Tasks



- Deploy a configuration directory to ZooKeeper
- Create a new collection
- Attach a local JConsole/VisualVM to a remote JVM
- Rolling restart (with Overseer awareness)
- Build Solr locally and patch remote
  - Use a relay server to scp the JARs to Amazon network once and then scp them to other nodes from within the network
- Put/get files
- Grep over all log files (across the cluster)

#### Other useful stuff ...



- fab mine: See clusters I'm running (or for other users too)
- fab kill\_mine: Terminate all instances I'm running
  - Use termination protection in production
- fab ssh\_to: Quick way to SSH to one of the nodes in a cluster
- fab stop/recover/kill: Basic commands for controlling specific Solr nodes in the cluster
- fab jmeter: Execute a JMeter test plan against your cluster
  - Example test plan and Java sampler is included with the source

## SolrCloud Tools (SolrJ client app)



# ./tools.sh -tool healthcheck

- Java-based command-line application that uses SolrJ's CloudSolrServer to perform advanced cluster management operations:
  - healthcheck: collect metadata and health information from all replicas for a collection from ZooKeeper
  - backup: create a snapshot of each shard in a collection for backing up to remote storage (S3)
- Framework for building complex tools that benefit from having access to cluster state information in ZooKeeper

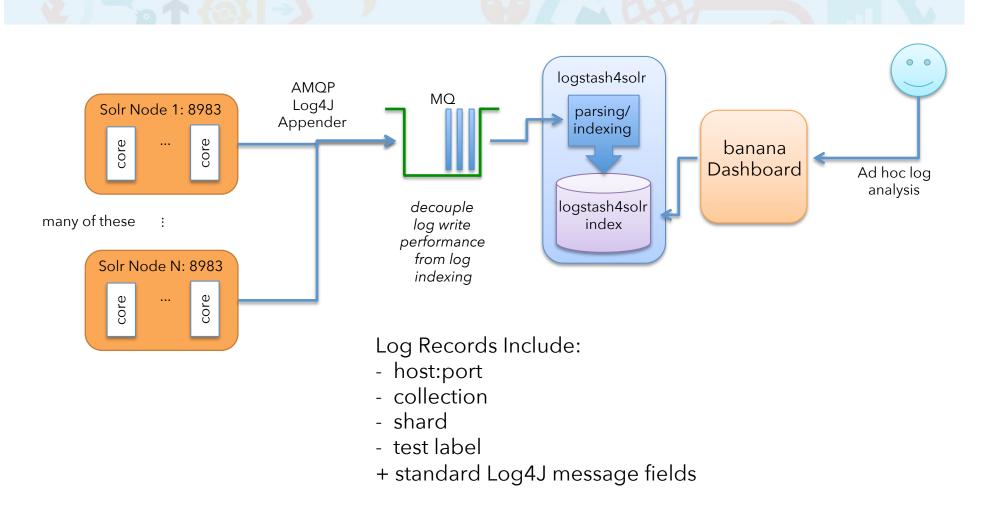
#### **SiLK Integration**



- SiLK: Solr integrated with Logstash and Kibana
  - Index time-series data, such as log data (collectd, Solr logs, ...)
  - Build cool dashboards with Banana (fork of Kibana)
- Easily aggregate all WARN and more severe log messages from all Solr servers into logstash4solr
- Send collectd metrics to logstash4solr

### **SiLK Integration**





#### What's Next?



- Migrate to using Apache libcloud instead of using boto directly
- Use this framework to perform large-scale performance testing
  - Report results back to community
- Ability to request spot instances
  - Good for testing only
- Chaos monkey tests
  - integrate jepsen?
- Open source so please kick the tires!

### Wrap-up



- LucidWorks: <a href="http://www.lucidworks.com">http://www.lucidworks.com</a>
- SiLK: <a href="http://www.lucidworks.com/lucidworks-silk/">http://www.lucidworks.com/lucidworks-silk/</a>
- Solr In Action: <a href="http://www.manning.com/grainger/">http://www.manning.com/grainger/</a>
- Connect: @thelabdude / tim.potter@lucidworks.com

#### **Questions?**

