Migrating High Performance Computing to the Cloud for a Bioinformatics Core - A Pilot Test

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#### What we do in computing

HPC Clusters

#### Storage Servers

#### Application Servers







#### > 50 labs/cores > 10 labs/cores



> 50 labs



## Computing challenges

- Storage capacity
- Long job queue
- Aging equipment
- Maintenance efforts
- No billing model





# **Cloud computing**

Instant Access

• Scalability



• Transparent cost



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# Why choose Amazon Web Services (AWS)

- MIT participates AWS service provided under Internet2
- MIT provides high-speed connection link to AWS





## Instant HPC: CfnCluster

- Cloud formation cluster (CfnCluster)
- Framework to build and manage High Performance Computing (HPC) clusters on AWS

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- Quick start in 10 min
- Elasticity using auto scaling











## IO performance test using fio



6 compute nodes (C4.2xlarge)





#### Performance test: bwa



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## **RNA-Seq Pipeline Test: rsem**



Total Project Cost on AWS: \$10





# **Genomics Core Illumina Pipeline**



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# Performance Test: Illumina Pipeline



## **Cost Estimation**

- On-premises HPC: \$20K-30K/year (not including MIT-provided infrastructure such as room, power, cooling and network)
- On-premises storage: \$50K/year
- Get wall clock from SGE history (\$qacct -d 365)
- AWS: 300,000 hr/year x \$0.4/hr = \$120K/year
- Parallel factor: assume 2-3 jobs running simultaneously on one node on average, HPC cost on AWS could be \$40K-60K/year
- Spot instances (saving cost but may lose productivity)
- AWS S3 storage cost: \$360/TB/year





# Thoughts on Cloud Computing

- Solve on-premises computing challenges (capacity planning, job queue, aging equipment, maintenance and billing)
- User training
- Need expert knowledge of AWS
- Hybrid environment





# Summary

- Cloud computing provided us expanded computing capacity
- AWS HPC performance comparable to our on-premises hardware



