

Genome Analysis in a Dynamically Scaled Hybrid Cloud

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In a Nutshell

- Users want to run analysis tasks
- Admin wants to dynamically allocate resources
- Non-expert users are bad at assigning resources to their jobs
- We propose to automatically control global and per-job parallelism...
- ...and show via simulation that this could work

Background: CELAR Project

- Cloud applications are often scalable
- Example: web database frontend
 - How many database servers?
 - How many web servers?
 - What spec cloud instance for each one?
- Parameters adjusted in ad-hoc manner, using admin's intuition
- CELAR: automate setting these parameters instead

Background: CELAR Project

- Applications expose *metrics*
 - Web example: avg. page load time, web server memory usage, ...
- Admin sets *goals*
 - Keep page load time below 1 second
- CELAR can hire and release cloud resources
- Learns how different resources help achieve the goals

Background: Resource Managers

- Users submit jobs
 - Annotated with *resource needs*
- Admin provides machines (usually fixed physical hardware, maybe cloud-hired)
- Resource manager (RM) queues jobs and assigns jobs to machines

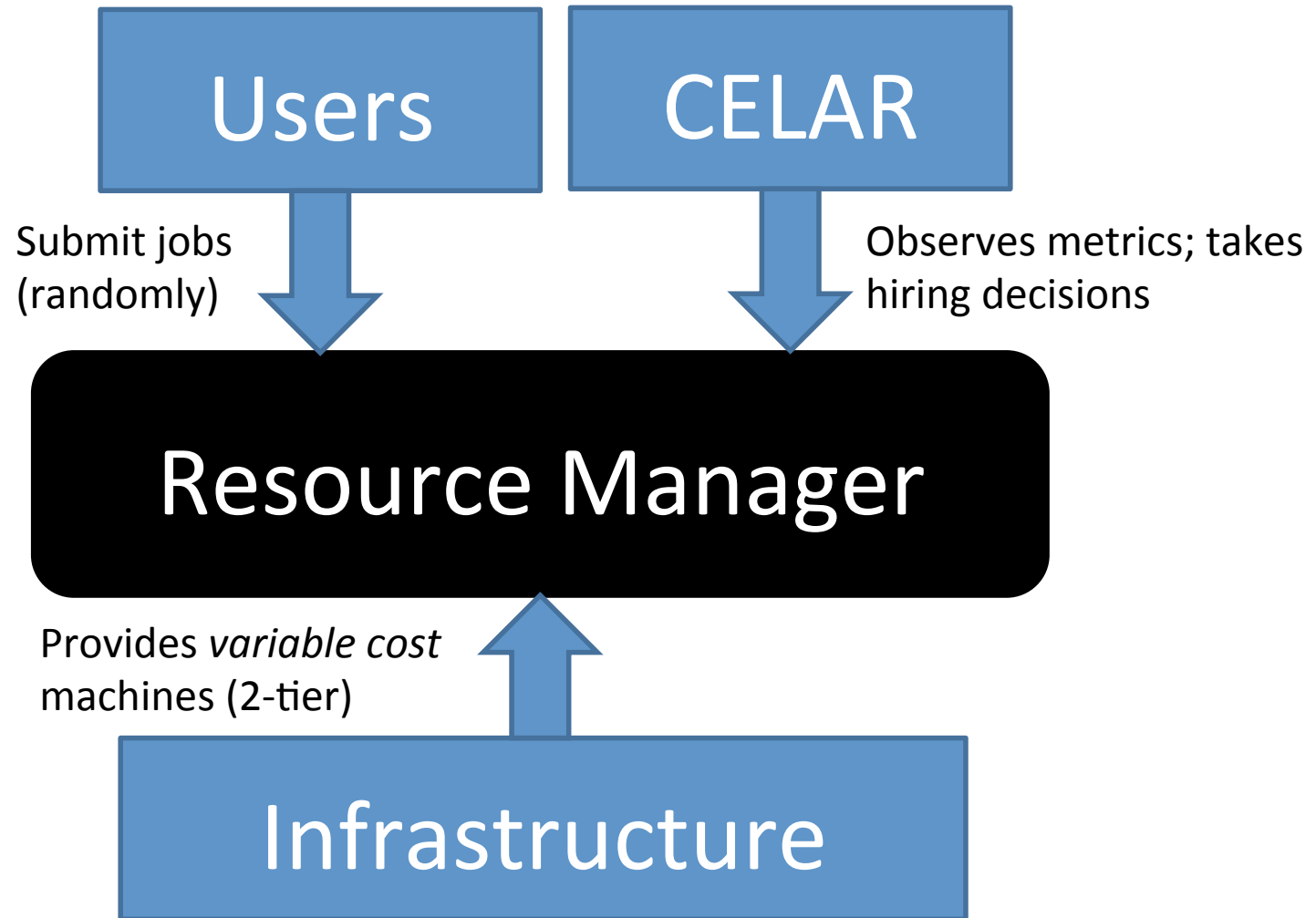
CELAR-ised Resource Manager

- Expose metrics to CELAR:
 - Job queue length, user “happiness”
- Set goals:
 - Maximise happiness!
- Let CELAR pick how many cloud VMs the RM gets, and what spec they have

This Work: Show that CELARising our RM can be a good thing

- CELAR chooses how many VMs the RM has at its disposal, and their spec
- Show that there is scope to improve over simple control policies
 - For example, hire a VM whenever there is a queue
- Evaluate by simulating interaction between users, resource manager and CELAR

Model



Model Users

- Users pay for service; we pay the infrastructure provider to hire machines
 - In practical academic environments, instead of money these might be *quota credits*
- Users pay more for “better” service
- We try to make as much “profit” as possible

Model Application: GATK

- 7-stage analysis pipeline
- Users care how long the *whole* pipeline takes
- Different stages have different scaling behaviour
 - Measured through profiling

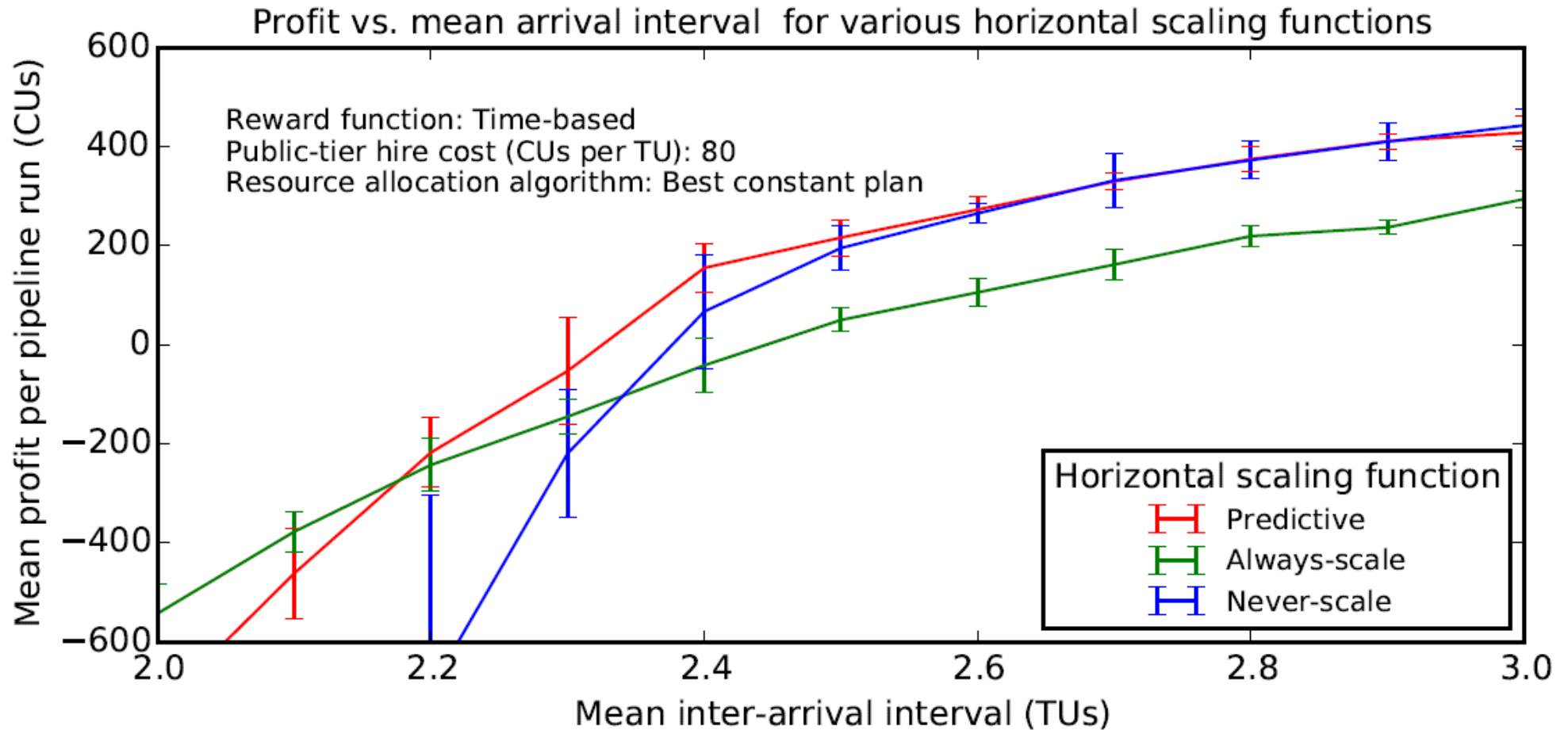
CELAR Task 1: When to hire, when to wait?

- Infrastructure has two cost tiers: cheap and expensive
- Suppose a queue is building up at the RM...
- ...but no cheap resources are available
- Should we hire an expensive machine or wait for a cheap one?

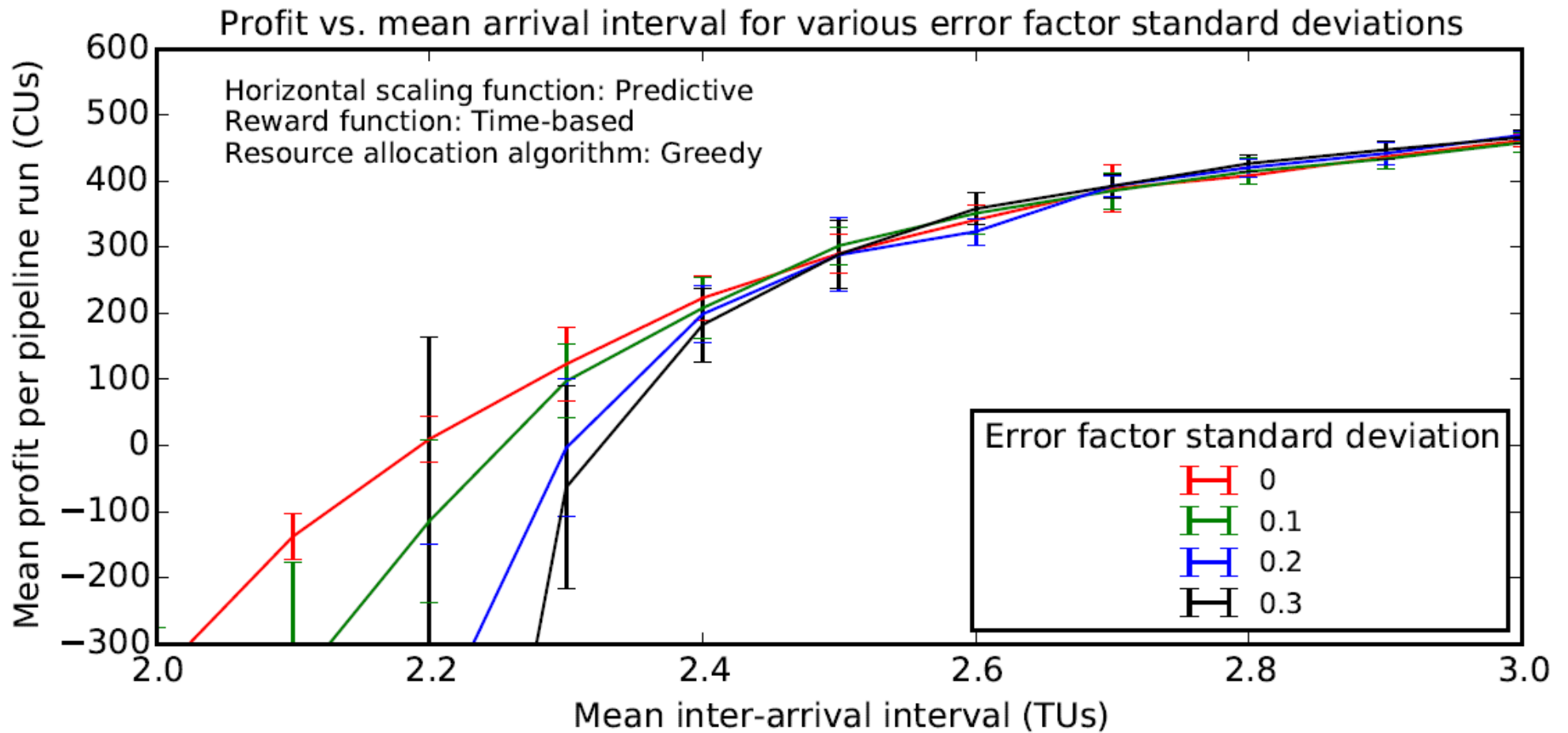
Proposed solution: Predict

- From our GATK profiling we can predict when running tasks will finish
- Weigh amount the user will pay for faster completion vs. expected extra cost
- If the queue is long, *all* queued jobs will benefit if we accelerate one

Predictive Scaling Example



Caveat: Prediction Quality



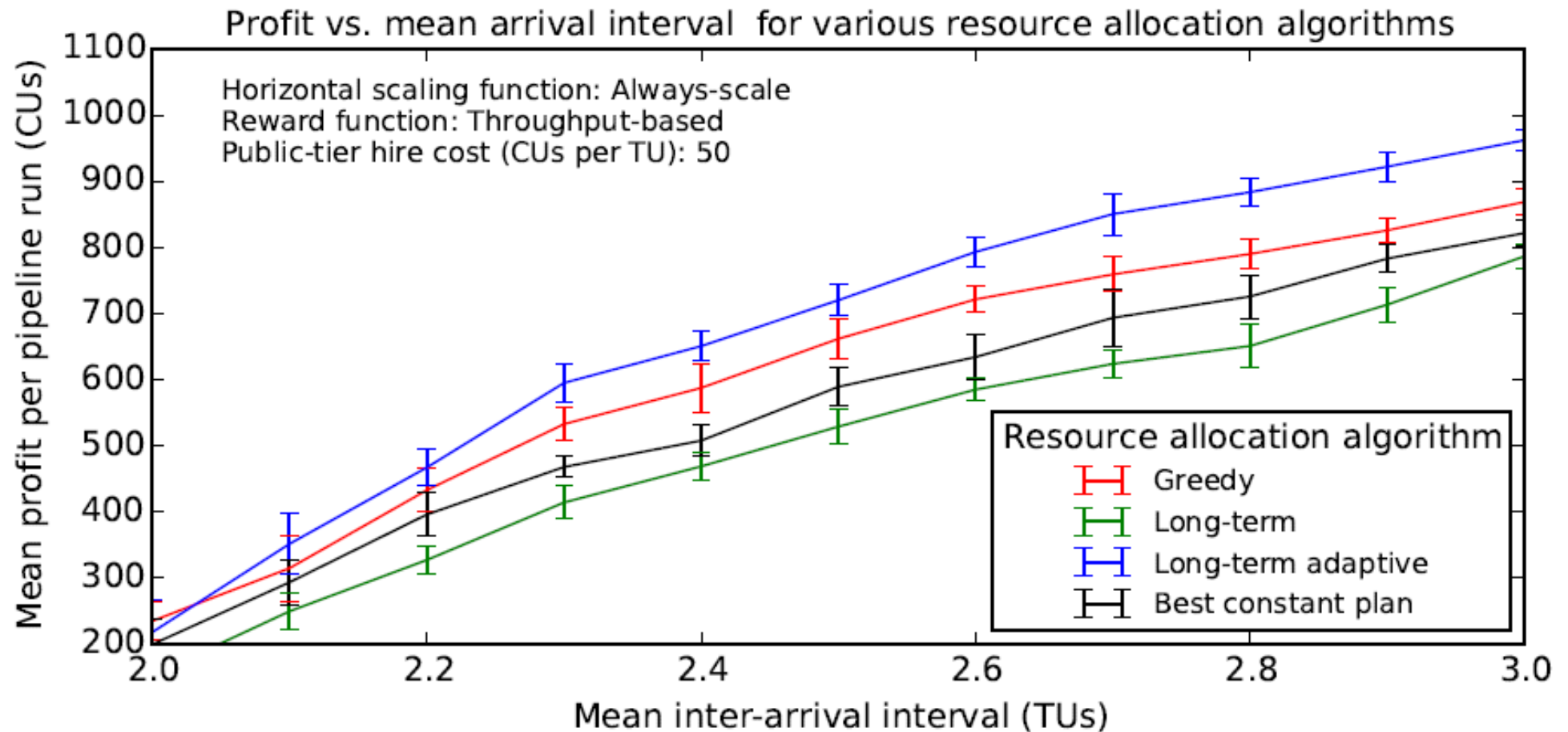
CELAR Task 2: To Thread or Not To Thread?

- Most GATK pipeline stages support multithreading...
- ...but some better than others, leading to diminishing returns
- During idle times, best to run multithreaded and make the users happy
- During busy times, best to run singlethreaded and maximise throughput

Proposed Solution: Weigh Expected Reward vs. Extra Cost

- Profiling lets us predict the effect of adding a thread
- Compare against cost, taking into account whether we would use cheap or expensive resources
- Three candidate schemes: greedy, long-term and long-term adaptive

Vertical Scaling Example



Caveat: Greed?

- If there are only 8 cheap cores left, and the next job would ideally use them all, should we really grab them all ourself?
- Surprisingly, yes!

Summary

- Developed candidate algorithms for automatically controlling the *size* of an RM's pool of worker VMs and the *parallelism* used by each job
- Showed ability to improve over baseline algorithms

What's Next?

- Apply findings in the real world: use auto-scaling on our real cluster
- Compare against existing work on e.g. Hadoop scheduling, where jobs are flexible by default
- Demonstrate within the CELAR framework

More Information

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