



# **Optimal Cluster and GPU Utilization with Run AI**

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# Optimal Cluster and GPU Utilization with Run:AI

The following sections provide details on the Run:AI installation, test scenarios, and results performed in this validation.

We validated the operation and performance of this system by using industry standard benchmark tools, including TensorFlow benchmarks. The ImageNet dataset was used to train ResNet-50, which is a famous Convolutional Neural Network (CNN) DL model for image classification. ResNet-50 delivers an accurate training result with a faster processing time, which enabled us to drive a sufficient demand on the storage.

[Next: Run AI Installation.](#)

## Run:AI Installation

To install Run:AI, complete the following steps:

1. Install the Kubernetes cluster using DeepOps and configure the NetApp default storage class.
2. Prepare GPU nodes:
  - a. Verify that NVIDIA drivers are installed on GPU nodes.
  - b. Verify that `nvidia-docker` is installed and configured as the default docker runtime.
3. Install Run:AI:
  - a. Log into the [Run:AI Admin UI](#) to create the cluster.
  - b. Download the created `runai-operator-<clustername>.yaml` file.
  - c. Apply the operator configuration to the Kubernetes cluster.

```
kubectl apply -f runai-operator-<clustername>.yaml
```

4. Verify the installation:
  - a. Go to <https://app.run.ai/>.
  - b. Go to the Overview dashboard.
  - c. Verify that the number of GPUs on the top right reflects the expected number of GPUs and the GPU nodes are all in the list of servers. For more information about Run:AI deployment, see [installing Run:AI on an on-premise Kubernetes cluster](#) and [installing the Run:AI CLI](#).

[Next: Run AI Dashboards and Views](#)

## Run:AI Dashboards and Views

After installing Run:AI on your Kubernetes cluster and configuring the containers correctly, you see the following dashboards and views on <https://app.run.ai> in your browser, as shown in the following figure.



There are 16 total GPUs in the cluster provided by two DGX-1 nodes. You can see the number of nodes, the total available GPUs, the allocated GPUs that are assigned with workloads, the total number of running jobs, pending jobs, and idle allocated GPUs. On the right side, the bar diagram shows GPUs per Project, which summarizes how different teams are using the cluster resource. In the middle is the list of currently running jobs with job details, including job name, project, user, job type, the node each job is running on, the number of GPU(s) allocated for that job, the current run time of the job, job progress in percentage, and the GPU utilization for that job. Note that the cluster is under-utilized (GPU utilization at 23%) because there are only three running jobs submitted by a single team (team-a).

In the following section, we show how to create multiple teams in the Projects tab and allocate GPUs for each team to maximize cluster usage and manage resources when there are many users per cluster. The test scenarios mimic enterprise environments in which memory and GPU resources are shared among training, inferencing, and interactive workloads.

[Next: Creating Projects for Data Science Teams and Allocating GPUs](#)

## Creating Projects for Data Science Teams and Allocating GPUs

Researchers can submit workloads through the Run:AI CLI, Kubeflow, or similar processes. To streamline resource allocation and create prioritization, Run:AI introduces the concept of Projects. Projects are quota entities that associate a project name with GPU allocation and preferences. It is a simple and convenient way to manage multiple data science teams.

A researcher submitting a workload must associate a project with a workload request. The Run:AI scheduler compares the request against the current allocations and the project and determines whether the workload can be allocated resources or whether it should remain in a pending state.

As a system administrator, you can set the following parameters in the Run:AI Projects tab:

- **Model projects.** Set a project per user, set a project per team of users, and set a project per a real organizational project.
- **Project quotas.** Each project is associated with a quota of GPUs that can be allocated for this project at the same time. This is a guaranteed quota in the sense that researchers using this project are guaranteed to get this number of GPUs no matter what the status in the cluster is. As a rule, the sum of the project allocation should be equal to the number of GPUs in the cluster. Beyond that, a user of this project can receive an over-quota. As long as GPUs are unused, a researcher using this project can get more GPUs. We demonstrate over-quota testing scenarios and fairness considerations in [Achieving High Cluster Utilization with Over-Quota GPU Allocation](#), [Basic Resource Allocation Fairness](#), and [Over-Quota Fairness](#).
- Create a new project, update an existing project, and delete an existing project.
- **Limit jobs to run on specific node groups.** You can assign specific projects to run only on specific nodes. This is useful when the project team needs specialized hardware, for example, with enough memory. Alternatively, a project team might be the owner of specific hardware that was acquired with a specialized budget, or when you might need to direct build or interactive workloads to work on weaker hardware and direct longer training or unattended workloads to faster nodes. For commands to group nodes and set affinity for a specific project, see the [Run:AI Documentation](#).
- **Limit the duration of interactive jobs.** Researchers frequently forget to close interactive jobs. This might lead to a waste of resources. Some organizations prefer to limit the duration of interactive jobs and close them automatically.

The following figure shows the Projects view with four teams created. Each team is assigned a different number of GPUs to account for different workloads, with the total number of GPUs equal to that of the total available GPUs in a cluster consisting of two DGX-1s.



| run:ai   Projects  |               | Cluster: cluster1   rick.huang@netapp.com |                        | NetApp                      |  |
|--|---------------|---|------------------------|-----------------------------|--|
| Filter and Search  |               | + Add New project                         |                        |                             |  |
| Project Name ↓   | Assigned GPUs | Created                                   | Training Node Affinity | Interactive Node Affinity ⚙ |  |
|  team-a | 2             | 07/27/20, 9:28AM                          | none                   | none                        |  |
|  team-b | 4             | 07/28/20, 7:50AM                          | none                   | none                        |  |
|  team-c | 2             | 07/28/20, 7:50AM                          | none                   | none                        |  |
|  team-d | 8             | 07/28/20, 7:51AM                          | none                   | none                        |  |

[Next: Submitting Jobs in Run AI CLI](#)

## Submitting Jobs in Run:AI CLI

This section provides the detail on basic Run:AI commands that you can use to run any Kubernetes job. It is divided into three parts according to workload type. AI/ML/DL workloads can be divided into two generic types:

- **Unattended training sessions.** With these types of workloads, the data scientist prepares a self-running workload and sends it for execution. During the execution, the customer can examine the results. This type of workload is often used in production or when model development is at a stage where no human intervention is required.

- **Interactive build sessions.** With these types of workloads, the data scientist opens an interactive session with Bash, Jupyter Notebook, remote PyCharm, or similar IDEs and accesses GPU resources directly. We include a third scenario for running interactive workloads with connected ports to reveal an internal port to the container user..

## Unattended Training Workloads

After setting up projects and allocating GPU(s), you can run any Kubernetes workload using the following command at the command line:

```
$ runai project set team-a runai submit hyper1 -i gcr.io/run-ai-  
demo/quickstart -g 1
```

This command starts an unattended training job for team-a with an allocation of a single GPU. The job is based on a sample docker image, `gcr.io/run-ai-demo/quickstart`. We named the job `hyper1`. You can then monitor the job's progress by running the following command:

```
$ runai list
```

The following figure shows the result of the `runai list` command. Typical statuses you might see include the following:

- **ContainerCreating.** The docker container is being downloaded from the cloud repository.
- **Pending.** The job is waiting to be scheduled.
- **Running.** The job is running.



```
~> runai list  
Showing jobs for project team-a  
NAME      STATUS  AGE  NODE                                     IMAGE                                     TYPE      PROJECT  USER  GPUs  
hyper1    Running  11s  gke-dev-yaron1-gpu-4-pool-154f511d-5nk5 gcr.io/run-ai-demo/quickstart          Train     team-a   yaron  1
```

To get an additional status on your job, run the following command:

```
$ runai get hyper1
```

To view the logs of the job, run the `runai logs <job-name>` command:

```
$ runai logs hyper1
```

In this example, you should see the log of a running DL session, including the current training epoch, ETA, loss function value, accuracy, and time elapsed for each step.

You can view the cluster status on the Run:AI UI at <https://app.run.ai/>. Under Dashboards > Overview, you can monitor GPU utilization.

To stop this workload, run the following command:

```
$ runai delte hyper1
```

This command stops the training workload. You can verify this action by running `runai list` again. For more detail, see [launching unattended training workloads](#).

## Interactive Build Workloads

After setting up projects and allocating GPU(s) you can run an interactive build workload using the following command at the command line:

```
$ runai submit build1 -i python -g 1 --interactive --command sleep --args infinity
```

The job is based on a sample docker image python. We named the job build1.



The `-- interactive` flag means that the job does not have a start or end. It is the researcher's responsibility to close the job. The administrator can define a time limit for interactive jobs after which they are terminated by the system.

The `--g 1` flag allocates a single GPU to this job. The command and argument provided is `--command sleep --args infinity`. You must provide a command, or the container starts and then exits immediately.

The following commands work similarly to the commands described in [Unattended Training Workloads](#):

- `runai list`: Shows the name, status, age, node, image, project, user, and GPUs for jobs.
- `runai get build1`: Displays additional status on the job build1.
- `runai delete build1`: Stops the interactive workload build1. To get a bash shell to the container, the following command:

```
$ runai bash build1
```

This provides a direct shell into the computer. Data scientists can then develop or finetune their models within the container.

You can view the cluster status on the Run:AI UI at <https://app.run.ai>. For more detail, see [starting and using interactive build workloads](#).

## Interactive Workloads with Connected Ports

As an extension of interactive build workloads, you can reveal internal ports to the container user when starting a container with the Run:AI CLI. This is useful for cloud environments, working with Jupyter Notebooks, or connecting to other microservices. [Ingress](#) allows access to Kubernetes services from outside the Kubernetes cluster. You can configure access by creating a collection of rules that define which inbound connections reach which services.

For better management of external access to the services in a cluster, we suggest that cluster administrators install [Ingress](#) and configure LoadBalancer.

To use Ingress as a service type, run the following command to set the method type and the ports when submitting your workload:

```
$ runai submit test-ingress -i jupyter/base-notebook -g 1 \
--interactive --service-type=ingress --port 8888 \
--args="--NotebookApp.base_url=test-ingress" --command=start-notebook.sh
```

After the container starts successfully, execute `runai list` to see the `SERVICE URL(S)` with which to access the Jupyter Notebook. The URL is composed of the ingress endpoint, the job name, and the port. For example, see <https://10.255.174.13/test-ingress-8888>.

For more details, see [launching an interactive build workload with connected ports](#).

Next: [Achieving High Cluster Utilization](#)

## Achieving High Cluster Utilization

In this section, we emulate a realistic scenario in which four data science teams each submit their own workloads to demonstrate the Run:AI orchestration solution that achieves high cluster utilization while maintaining prioritization and balancing GPU resources. We start by using the ResNet-50 benchmark described in the section [ResNet-50 with ImageNet Dataset Benchmark Summary](#):

```
$ runai submit netapp1 -i netapp/tensorflow-tf1-py3:20.01.0 --local-image
--large-shm -v /mnt:/mnt -v /tmp:/tmp --command python --args
"/netapp/scripts/run.py" --args "--
dataset_dir=/mnt/mount_0/dataset/imagenet/imagenet_original/" --args "--
num_mounts=2" --args "--dgx_version=dgx1" --args "--num_devices=1" -g 1
```

We ran the same ResNet-50 benchmark as in [NVA-1121](#). We used the flag `--local-image` for containers not residing in the public docker repository. We mounted the directories `/mnt` and `/tmp` on the host DGX-1 node to `/mnt` and `/tmp` to the container, respectively. The dataset is at NetApp AFFA800 with the `dataset_dir` argument pointing to the directory. Both `--num_devices=1` and `-g 1` mean that we allocate one GPU for this job. The former is an argument for the `run.py` script, while the latter is a flag for the `runai submit` command.

The following figure shows a system overview dashboard with 97% GPU utilization and all sixteen available GPUs allocated. You can easily see how many GPUs are allocated for each team in the GPUs/Project bar chart. The Running Jobs pane shows the current running job names, project, user, type, node, GPUs consumed, run time, progress, and utilization details. A list of workloads in queue with their wait time is shown in Pending Jobs. Finally, the Nodes box offers GPU numbers and utilization for individual DGX-1 nodes in the cluster.





Next: [Fractional GPU Allocation for Less Demanding or Interactive Workloads](#)

## Fractional GPU Allocation for Less Demanding or Interactive Workloads

When researchers and developers are working on their models, whether in the development, hyperparameter tuning, or debugging stages, such workloads usually require fewer computational resources. It is therefore more efficient to provision fractional GPU and memory such that the same GPU can simultaneously be allocated to other workloads. Run:AI's orchestration solution provides a fractional GPU sharing system for containerized workloads on Kubernetes. The system supports workloads running CUDA programs and is especially suited for lightweight AI tasks such as inference and model building. The fractional GPU system transparently gives data science and AI engineering teams the ability to run multiple workloads simultaneously on a single GPU. This enables companies to run more workloads, such as computer vision, voice recognition, and natural language processing on the same hardware, thus lowering costs.

Run:AI's fractional GPU system effectively creates virtualized logical GPUs with their own memory and computing space that containers can use and access as if they were self-contained processors. This enables several workloads to run in containers side-by-side on the same GPU without interfering with each other. The solution is transparent, simple, and portable and it requires no changes to the containers themselves.

A typical usecase could see two to eight jobs running on the same GPU, meaning that you could do eight times the work with the same hardware.

For the job `frac05` belonging to project `team-d` in the following figure, we can see that the number of GPUs allocated was 0.50. This is further verified by the `nvidia-smi` command, which shows that the GPU memory available to the container was 16,255MB: half of the 32GB per V100 GPU in the DGX-1 node.

```
root@run-deploy:~# runai bash frac05 -p team-d
root@frac05-0:/workload# nvidia-smi
Tue Jul 28 15:17:03 2020

+-----+
| NVIDIA-SMI 450.51.05      Driver Version: 450.51.05      CUDA Version: 11.0      |
+-----+-----+
| GPU   Name                Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC |
| Fan   Temp   Perf    Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M. |
|                               |                    |            MIG M. |
+-----+-----+
|  0  Tesla V100-SXM2...    On         | 00000000:07:00.0 Off |                    0 |
| N/A   57C    P0     240W / 300W | 15525MiB / 16255MiB |   100%    Default  |
|                               |                    |            N/A   |
+-----+-----+

+-----+
| Processes: |
| GPU   GI    CI          PID    Type    Process name          GPU Memory |
|                               |                      |              Usage |
+-----+-----+
|  0   N/A   N/A         156      C      python3              15525MiB |
+-----+-----+
```

Next: [Achieving High Cluster Utilization with Over-Quota GPU Allocation](#)

## Achieving High Cluster Utilization with Over-Quota GPU Allocation

In this section and in the sections [Basic Resource Allocation Fairness](#), and [Over-Quota Fairness](#), we have devised advanced testing scenarios to demonstrate the Run:AI orchestration capabilities for complex workload management, automatic preemptive scheduling, and over-quota GPU provisioning. We did this to achieve high cluster-resource usage and optimize enterprise-level data science team productivity in an ONTAP AI environment.

For these three sections, set the following projects and quotas:

| Project | Quota |
|---------|-------|
| team-a  | 4     |
| team-b  | 2     |
| team-c  | 2     |

| Project | Quota |
|---------|-------|
| team-d  | 8     |

In addition, we use the following containers for these three sections:

- Jupyter Notebook: `jupyter/base-notebook`
- Run:AI quickstart: `gcr.io/run-ai-demo/quickstart`

We set the following goals for this test scenario:

- Show the simplicity of resource provisioning and how resources are abstracted from users
- Show how users can easily provision fractions of a GPU and integer number of GPUs
- Show how the system eliminates compute bottlenecks by allowing teams or users to go over their resource quota if there are free GPUs in the cluster
- Show how data pipeline bottlenecks are eliminated by using the NetApp solution when running compute-intensive jobs, such as the NetApp container
- Show how multiple types of containers are running using the system
  - Jupyter Notebook
  - Run:AI container
- Show high utilization when the cluster is full

For details on the actual command sequence executed during the testing, see [Testing Details for Section 4.8](#).

When all 13 workloads are submitted, you can see a list of container names and GPUs allocated, as shown in the following figure. We have seven training and six interactive jobs, simulating four data science teams, each with their own models running or in development. For interactive jobs, individual developers are using Jupyter Notebooks to write or debug their code. Thus, it is suitable to provision GPU fractions without using too many cluster resources.

```
root@run-deploy:~# runai list -A
```

| NAME          | STATUS  | AGE | NODE   | IMAGE                         | TYPE        | PROJECT | USER | GPUs | CREATED BY | CLI | SERVICE URL(S)   |
|---------------|---------|-----|--------|-------------------------------|-------------|---------|------|------|------------|-----|--|
| b-4-gg        | Running | 2m  | dgx1-2 | gcr.io/run-ai-demo/quickstart | Train       | team-b  | root | 2    | true       |     |  |
| c-5-g         | Running | 2m  | dgx1-2 | gcr.io/run-ai-demo/quickstart | Train       | team-c  | root | 1    | true       |     |  |
| c-4-gg        | Running | 2m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Train       | team-c  | root | 2    | true       |     |  |
| b-3-g         | Running | 2m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Train       | team-b  | root | 1    | true       |     |  |
| c-3-g02       | Running | 2m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Interactive | team-c  | root | 0.2  | true       |     |  |
| d-1-gggg      | Running | 2m  | dgx1-2 | gcr.io/run-ai-demo/quickstart | Train       | team-d  | root | 4    | true       |     |  |
| c-2-g03       | Running | 2m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Interactive | team-c  | root | 0.3  | true       |     |  |
| c-1-g05       | Running | 2m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Interactive | team-c  | root | 0.5  | true       |     |  |
| a-2-gg        | Running | 3m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Train       | team-a  | root | 2    | true       |     |  |
| b-2-g04       | Running | 3m  | dgx1-2 | gcr.io/run-ai-demo/quickstart | Interactive | team-b  | root | 0.4  | true       |     |  |
| a-1-g         | Running | 3m  | dgx1-1 | gcr.io/run-ai-demo/quickstart | Train       | team-a  | root | 1    | true       |     |  |
| b-1-g06       | Running | 3m  | dgx1-2 | gcr.io/run-ai-demo/quickstart | Interactive | team-b  | root | 0.6  | true       |     |  |
| a-1-1-jupyter | Running | 3m  | dgx1-1 | jupyter/base-notebook         | Interactive | team-a  | root | 1    | true       |     | <a href="http://10.61.218.134/a-1-1-jupyter">http://10.61.218.134/a-1-1-jupyter</a> ,<br><a href="https://10.61.218.134/a-1-1-jupyter">https://10.61.218.134/a-1-1-jupyter</a> |

The results of this testing scenario show the following:

- The cluster should be full: 16/16 GPUs are used.
- High cluster utilization.
- More experiments than GPUs due to fractional allocation.
- team-d is not using all their quota; therefore, team-b and team-c can use additional GPUs for their experiments, leading to faster time to innovation.

# Basic Resource Allocation Fairness

In this section, we show that, when `team-d` asks for more GPUs (they are under their quota), the system pauses the workloads of `team-b` and `team-c` and moves them into a pending state in a fair-share manner.

For details including job submissions, container images used, and command sequences executed, see the section [Testing Details for Section 4.9](#).

The following figure shows the resulting cluster utilization, GPUs allocated per team, and pending jobs due to automatic load balancing and preemptive scheduling. We can observe that when the total number of GPUs requested by all team workloads exceeds the total available GPUs in the cluster, Run:AI's internal fairness algorithm pauses one job each for `team-b` and `team-c` because they have met their project quota. This provides overall high cluster utilization while data science teams still work under resource constraints set by an administrator.



The results of this testing scenario demonstrate the following:

- **Automatic load balancing.** The system automatically balances the quota of the GPUs, such that each team is now using their quota. The workloads that were paused belong to teams that were over their quota.
- **Fair share pause.** The system chooses to stop the workload of one team that was over their quota and then stop the workload of the other team. Run:AI has internal fairness algorithms.

# Over-Quota Fairness

In this section, we expand the scenario in which multiple teams submit workloads and exceed their quota. In this way, we demonstrate how Run:AI's fairness algorithm allocates cluster resources according to the ratio of preset quotas.

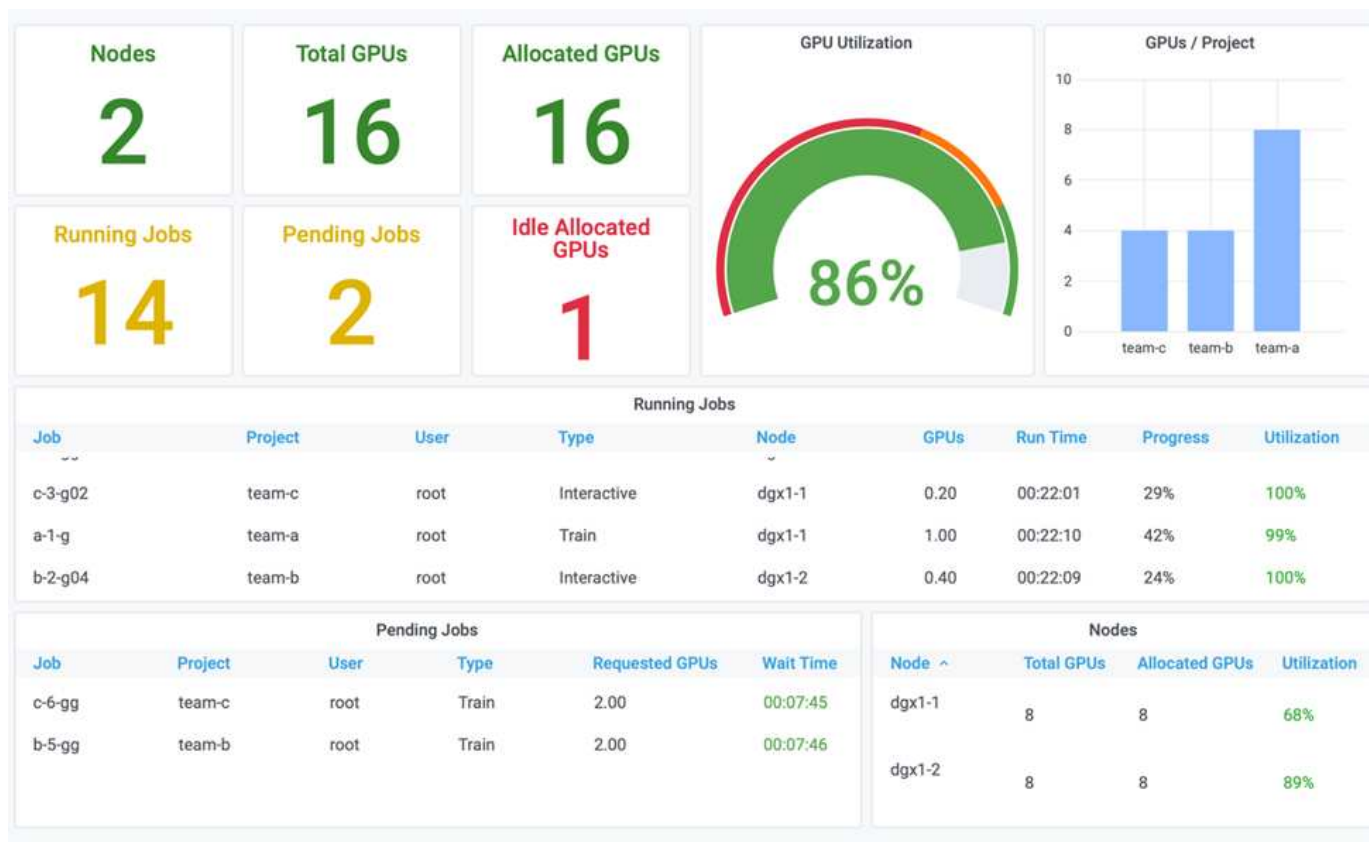
Goals for this test scenario:

- Show queuing mechanism when multiple teams are requesting GPUs over their quota.
- Show how the system distributes a fair share of the cluster between multiple teams that are over their quota according to the ratio between their quotas, so that the team with the larger quota gets a larger share of the spare capacity.

At the end of [Basic Resource Allocation Fairness](#), there are two workloads queued: one for team-b and one for team-c. In this section, we queue additional workloads.

For details including job submissions, container images used, and command sequences executed, see [Testing Details for section 4.10](#).

When all jobs are submitted according to the section [Testing Details for section 4.10](#), the system dashboard shows that team-a, team-b, and team-c all have more GPUs than their preset quota. team-a occupies four more GPUs than its preset soft quota (four), whereas team-b and team-c each occupy two more GPUs than their soft quota (two). The ratio of over-quota GPUs allocated is equal to that of their preset quota. This is because the system used the preset quota as a reference of priority and provisioned accordingly when multiple teams request more GPUs, exceeding their quota. Such automatic load balancing provides fairness and prioritization when enterprise data science teams are actively engaged in AI model development and production.





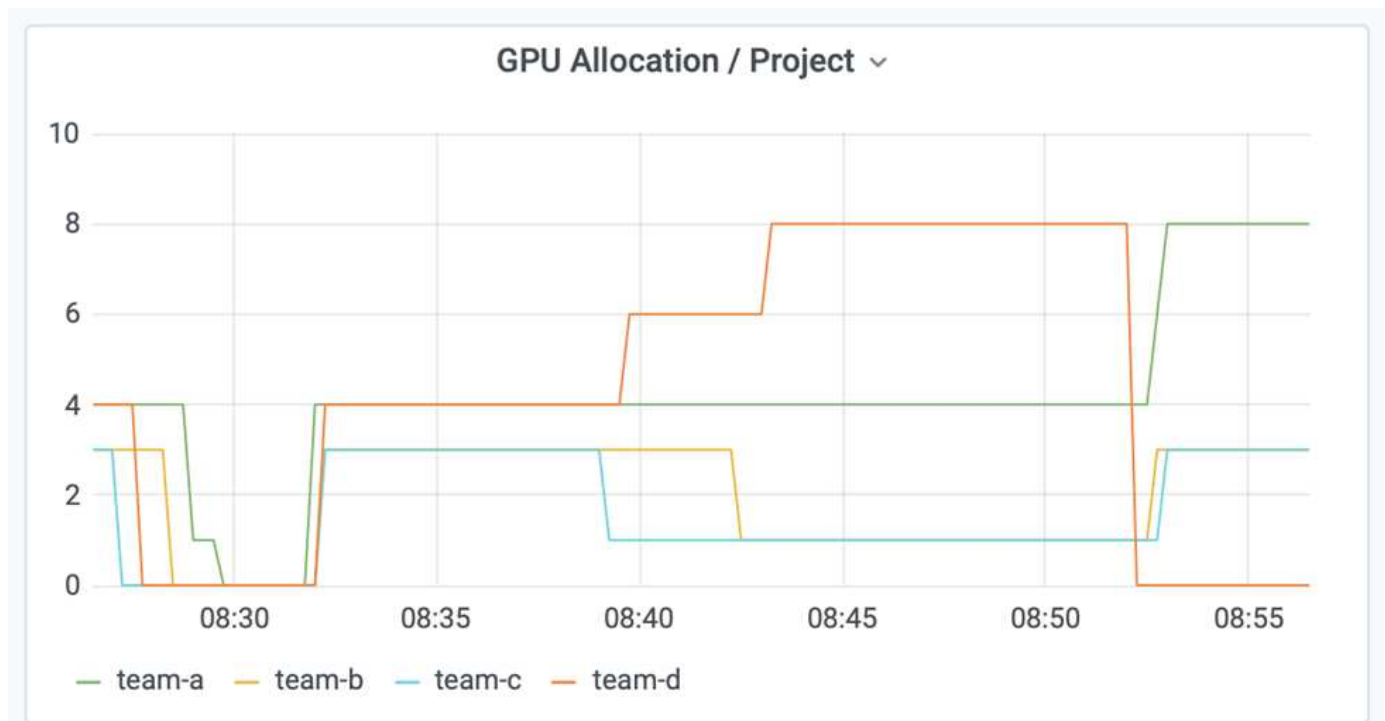
The results of this testing scenario show the following:

- The system starts to de-queue the workloads of other teams.
- The order of the dequeuing is decided according to fairness algorithms, such that `team-b` and `team-c` get the same amount of over-quota GPUs (since they have a similar quota), and `team-a` gets a double amount of GPUs since their quota is two times higher than the quota of `team-b` and `team-c`.
- All the allocation is done automatically.

Therefore, the system should stabilize on the following states:

| Project | GPUs allocated | Comment                                       |
|---------|----------------|---|
| team-a  | 8/4            | Four GPUs over the quota. Empty queue.        |
| team-b  | 4/2            | Two GPUs over the quota. One workload queued. |
| team-c  | 4/2            | Two GPUs over the quota. One workload queued. |
| team-d  | 0/8            | Not using GPUs at all, no queued workloads.   |

The following figure shows the GPU allocation per project over time in the Run:AI Analytics dashboard for the sections [Achieving High Cluster Utilization with Over-Quota GPU Allocation](#), [Basic Resource Allocation Fairness](#), and [Over-Quota Fairness](#). Each line in the figure indicates the number of GPUs provisioned for a given data science team at any time. We can see that the system dynamically allocates GPUs according to workloads submitted. This allows teams to go over quota when there are available GPUs in the cluster, and then preempt jobs according to fairness, before finally reaching a stable state for all four teams.



[Next: Saving Data to a Trident-Provisioned PersistentVolume](#)

# Saving Data to a Trident-Provisioned PersistentVolume

NetApp Trident is a fully supported open source project designed to help you meet the sophisticated persistence demands of your containerized applications. You can read and write data to a Trident-provisioned Kubernetes PersistentVolume (PV) with the added benefit of data tiering, encryption, NetApp Snapshot technology, compliance, and high performance offered by NetApp ONTAP data management software.

## Reusing PVCs in an Existing Namespace

For larger AI projects, it might be more efficient for different containers to read and write data to the same Kubernetes PV. To reuse a Kubernetes Persistent Volume Claim (PVC), the user must have already created a PVC. See the [NetApp Trident documentation](#) for details on creating a PVC. Here is an example of reusing an existing PVC:

```
$ runai submit pvc-test -p team-a --pvc test:/tmp/pvc1mount -i gcr.io/run-ai-demo/quickstart -g 1
```

Run the following command to see the status of job `pvc-test` for project `team-a`:

```
$ runai get pvc-test -p team-a
```

You should see the PV `/tmp/pvc1mount` mounted to `team-a` job `pvc-test`. In this way, multiple containers can read from the same volume, which is useful when there are multiple competing models in development or in production. Data scientists can build an ensemble of models and then combine prediction results by majority voting or other techniques.

Use the following to access the container shell:

```
$ runai bash pvc-test -p team-a
```

You can then check the mounted volume and access your data within the container.

This capability of reusing PVCs works with NetApp FlexVol volumes and NetApp ONTAP FlexGroup volumes, enabling data engineers more flexible and robust data management options to leverage your data fabric powered by NetApp.

[Next: Conclusion](#)

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