



# **Example Kubeflow Operations and Tasks**

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# Example Kubeflow Operations and Tasks

This section includes examples of various operations and tasks that you may want to perform using Kubeflow.

Next: [Provision a Jupyter Notebook Workspace for Data Scientist or Developer Use.](#)

## Example Kubeflow Operations and Tasks

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Next: [Provision a Jupyter Notebook Workspace for Data Scientist or Developer Use.](#)

## Provision a Jupyter Notebook Workspace for Data Scientist or Developer Use

Kubeflow is capable of rapidly provisioning new Jupyter Notebook servers to act as data scientist workspaces. To provision a new Jupyter Notebook server with Kubeflow, perform the following tasks. For more information about Jupyter Notebooks within the Kubeflow context, see the [official Kubeflow documentation](#).

1. From the Kubeflow central dashboard, click Notebook Servers in the main menu to navigate to the Jupyter Notebook server administration page.



2. Click New Server to provision a new Jupyter Notebook server.



3. Give your new server a name, choose the Docker image that you want your server to be based on, and specify the amount of CPU and RAM to be reserved by your server. If the Namespace field is blank, use the Select Namespace menu in the page header to choose a namespace. The Namespace field is then auto-populated with the chosen namespace.

In the following example, the `kubeflow-anonymous` namespace is chosen. In addition, the default values for Docker image, CPU, and RAM are accepted.



The screenshot shows the Kubeflow Central Dashboard in a web browser. The browser's address bar shows the URL `10.61.218.131:31380/_jupyter/?ns=kubeflow-anonym...`. The dashboard header includes the Kubeflow logo and a dropdown menu for the namespace, which is currently set to `kubeflow-anonymous`. The main content area is titled "Name" and contains a form for specifying the Notebook Server details. The form has three sections: "Name", "Image", and "CPU / RAM".

**Name**  
Specify the name of the Notebook Server and the Namespace it will belong to.

Name	Namespace
mike	kubeflow-anonymous

**Image**  
A starter Jupyter Docker Image with a baseline deployment and typical ML packages.

☐ Custom Image

Image: `gcr.io/kubeflow-images-public/tensorflow-1.13.1-notebook-cpu:v0.5.0`

**CPU / RAM**  
Specify the total amount of CPU and RAM reserved by your Notebook Server. For CPU-intensive workloads, you can choose more than 1 CPU (e.g. 1.5).

CPU	Memory
0.5	1.0Gi

4. Specify the workspace volume details. If you choose to create a new volume, then that volume or PVC is provisioned using the default StorageClass. Because a StorageClass utilizing Trident was designated as the default StorageClass in the section [Kubeflow Deployment](#), the volume or PVC is provisioned with Trident. This volume is automatically mounted as the default workspace within the Jupyter Notebook Server container. Any notebooks that a user creates on the server that are not saved to a separate data volume are automatically saved to this workspace volume. Therefore, the notebooks are persistent across reboots.

**Workspace Volume**

Configure the Volume to be mounted as your personal Workspace.

☐ Don't use Persistent Storage for User's home

Type	Name	Size	Mode	Mount Point
New	workspace-mike	10Gi	ReadWriteOnce	/home/jovyan

- Add data volumes. The following example specifies an existing PVC named 'pb-fg-all' and accepts the default mount point.

**Data Volumes**

Configure the Volumes to be mounted as your Datasets.

[+ ADD VOLUME](#)

Type	Name	Size	Mode	Mount Point
Existing	pb-fg-all	10Gi	ReadWriteOnce	/home/jovyan/data-vol-1

- Optional:** Request that the desired number of GPUs be allocated to your notebook server. In the following example, one GPU is requested.

**Configurations**

Extra layers of configurations that will be applied to the new Notebook. (e.g. Insert credentials as Secrets, set Environment Variables.)

Configurations

**Extra Resources**

Specify extra resources that might be needed in the Notebook Server.

☒ Enable Shared Memory

Extra Resources \*

`{"nvidia.com/gpu": 1}`

Extra Resources available in the cluster (ex. NVIDIA GPUs)

**LAUNCH** **CANCEL**

- Click Launch to provision your new notebook server.
- Wait for your notebook server to be fully provisioned. This can take several minutes if you have never provisioned a server using the Docker image that you specified because the image needs to be downloaded. When your server has been fully provisioned, you see a green check mark in the Status column on the Jupyter Notebook server administration page.

**Notebook Servers** [+ NEW SERVER](#)

Status	Name	Age	Image	CPU	Memory	Volumes	
	mike	12 mins ago	tensorflow-1.13.1-notebook-cpu:v0.5.0	0.5	1.0Gi		<a href="#">CONNECT</a>

9. Click Connect to connect to your new server web interface.
10. Confirm that the dataset volume that was specified in step 6 is mounted on the server. Note that this volume is mounted within the default workspace by default. From the perspective of the user, this is just another folder within the workspace. The user, who is likely a data scientist and not an infrastructure expert, does not need to possess any storage expertise in order to use this volume.

**jupyter** [Quit](#)

Files Running Clusters

Select items to perform actions on them. [Upload](#) [New](#) [Refresh](#)

	Name	Last Modified	File size
<input type="checkbox"/>	/		
<input type="checkbox"/>	data-vol-1	a day ago	



- Open a Terminal and, assuming that a new volume was requested in step 5, execute `df -h` to confirm that a new Trident-provisioned persistent volume is mounted as the default workspace.

The default workspace directory is the base directory that you are presented with when you first access the server's web interface. Therefore, any artifacts that you create by using the web interface are stored on this Trident-provisioned persistent volume.





```

$ df -h
Filesystem                                Size  Used Avail
Use% Mounted on
overlay                                  439G   34G  382G
9% /
tmpfs                                     64M    0   64M
0% /dev
tmpfs                                    252G    0  252G
0% /sys/fs/cgroup
/dev/sda2                                439G   34G  382G
9% /etc/hosts
192.168.11.11:/trident_pvc_3dcfe7e5_d5a9_11e9_9b9d_00505681a82d 10G  320K   10G
1% /home/jovyan
tmpfs                                    252G    0  252G
0% /dev/shm
192.168.11.11:/pb_fg_all                  10T   10T   47G
100% /home/jovyan/data-vol-1
tmpfs                                    252G   12K  252G
1% /run/secrets/kubernetes.io/serviceaccount
tmpfs                                    252G   12K  252G
1% /proc/driver/nvidia
tmpfs                                    51G   4.9M   51G
1% /run/nvidia-persistenced/socket
udev                                     252G    0  252G
0% /dev/nvidia5
tmpfs                                    252G    0  252G
0% /proc/acpi
tmpfs                                    252G    0  252G
0% /proc/scsi
tmpfs                                    252G    0  252G
0% /sys/firmware
$

```

- Using the terminal, run `nvidia-smi` to confirm that the correct number of GPUs were allocated to the notebook server. In the following example, one GPU has been allocated to the notebook server as requested in step 7.

```

$ nvidia-smi
Fri Sep 13 13:52:15 2019
+-----+
| NVIDIA-SMI 410.104      | Driver Version: 410.104      | CUDA Version: N/A      |
+-----+-----+
| GPU  Name                Persistence-M| Bus-Id        Disp.A | Volatile Uncorr. ECC  |
| Fan  Temp  Perf  Pwr:Usage/Cap|      Memory-Usage | GPU-Util  Compute M.  |
+-----+-----+
| 0   Tesla V100-SXM2...    On         | 00000000:86:00:0 Off  |          0%          0 |
| N/A   38C    P0     46W / 300W | 0MiB / 32480MiB |              Default  |
+-----+-----+

+-----+
| Processes:                 GPU Memory |
| GPU       PID    Type    Process name                     Usage |
+-----+-----+
| No running processes found |
+-----+
$

```

Next: [Example Notebooks and Pipelines.](#)

## Example Notebooks and Pipelines

The [NetApp Data Science Toolkit for Kubernetes](#) can be used in conjunction with Kubeflow. Using the NetApp Data Science Toolkit with Kubeflow provides the following benefits:

- Data scientists can perform advanced NetApp data management operations directly from within a Jupyter Notebook.
- Advanced NetApp data management operations can be incorporated into automated workflows using the Kubeflow Pipelines framework.

Refer to the [Kubeflow Examples](#) section within the NetApp Data Science Toolkit GitHub repository for details on using the toolkit with Kubeflow.

[Next: Apache Airflow Deployment.](#)

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