ML Pipeline with Kubeflow on Azure: A Comprehensive Guide

Comprehensive Guide: ML Pipeline with Kubeflow on Azure

Prerequisites

1. System Requirements

- Linux/macOS/Windows with WSL2
- Python 3.8+
- · Docker Desktop
- Git

2. Cloud & Tools

```
# Install Azure CLI
curl -sL <a href="https://aka.ms/InstallAzureCLIDeb"> | sudo bash

# Install kubectl
curl -LO "<a href="https://dl.k8s.io/release/$(curl> -L -s"> <a href="https://dl.k8s.io/release/$(curl> -L -s"> <a href="https://dl.k8s.io/release/$table.txt"> + L -s"> <a href="https://dl.k8s.io/release/$table.txt"> + L -s"> <a href="https://dl.k8s.io/release/$table.txt</a>)/bin/linux/amd64/kubectl>"
sudo install -o root -g root -m 0755 kubectl /usr/local/bin/kubectl

# Install Kustomize
curl -s "<a href="https://raw.githubusercontent.com/kubernetes-sigs/kustomize/master/hack/install_kustomize.sh"> | bash
sudo mv kustomize /usr/local/bin/
# Python packages
pip install azure-cli azure-ml-sdk kubernetes kfp azureml-sdk pandas scikit-learn tensorflow</a>
```

3. Azure Resources

- Active Azure subscription
- · Resource group
- Storage account
- Azure Container Registry (ACR)

Setup Infrastructure

1. Create AKS Cluster

```
1 # Create resource group
2 az group create --name ml-resources --location eastus
3
4 # Create AKS cluster
5 az aks create \
6 --resource-group ml-resources \
7 --name ml-cluster \
8 --node-count 3 \
9 --enable-addons monitoring \
10 --generate-ssh-keys \
```

```
--node-vm-size Standard_DS3_v2

13  # Get credentials

14 az aks get-credentials --resource-group ml-resources --name ml-cluster
```

2. Deploy Kubeflow

```
# Clone manifests
git clone <a href="https://github.com/kubeflow/manifests.git">https://github.com/kubeflow/manifests.git</a>
cd manifests

# Install Kubeflow
while ! kustomize build example | kubectl apply -f -; do
echo "Retrying to apply resources";
sleep 30;
done

# Verify deployment
kubectl get pods -n kubeflow
```

Sample ML Pipeline Implementation

1. Dataset Preparation

```
1 # dataset prep.py
2 import pandas as pd
3 from sklearn.datasets import load_diabetes
4 from sklearn.model_selection import train_test_split
6 def prepare_data():
7
     # Load sample diabetes dataset
8
     diabetes = load_diabetes()
     df = pd.DataFrame(diabetes.data, columns=diabetes.feature_names)
     df['target'] = diabetes.target
10
11
13
     train_df, test_df = train_test_split(df, test_size=0.2, random_state=42)
14
     # Save to files
15
16
     train_df.to_csv('train.csv', index=False)
17
     test_df.to_csv('test.csv', index=False)
18
19
      return 'train.csv', 'test.csv'
20
21 if __name__ == '__main__':
      prepare_data()
```

2. Model Training

```
1 # train.py
2 import pandas as pd
3 import numpy as np
4 from sklearn.linear_model import Ridge
5 import pickle
6
7 def train_model(data_path):
8 # Load training data
```

```
train data = pd.read csv(data path)
10
       X = train_data.drop('target', axis=1)
       y = train_data['target']
11
12
13
       # Train model
14
       model = Ridge(alpha=1.0)
15
       model.fit(X, y)
16
17
       # Save model
       with open('model.pkl', 'wb') as f:
18
19
           pickle.dump(model, f)
20
       return 'model.pkl'
21
22
23 if __name__ == '__main__':
24
      import argparse
25
       parser = argparse.ArgumentParser()
26
       parser.add_argument('--data', type=str)
27
       args = parser.parse_args()
28
       train_model(args.data)
```

3. Model Evaluation

```
1 # evaluate.py
2 import pandas as pd
3 import pickle
4 from sklearn.metrics import mean_squared_error, r2_score
5 import json
6
7 def evaluate_model(model_path, test_data_path):
8
       # Load model and test data
9
       with open(model_path, 'rb') as f:
10
           model = pickle.load(f)
11
12
       test_data = pd.read_csv(test_data_path)
13
     X_test = test_data.drop('target', axis=1)
14
       y test = test data['target']
15
16
       # Make predictions
17
       y_pred = model.predict(X_test)
18
19
       # Calculate metrics
20
       metrics = {
21
           'mse': mean_squared_error(y_test, y_pred),
22
           'r2': r2_score(y_test, y_pred)
23
       }
24
25
       # Save metrics
       with open('metrics.json', 'w') as f:
26
27
           json.dump(metrics, f)
28
29
       return 'metrics.json'
30
31 if __name__ == '__main__':
32
      import argparse
33
       parser = argparse.ArgumentParser()
34
       parser.add_argument('--model', type=str)
35
       parser.add_argument('--test-data', type=str)
```

```
args = parser.parse_args()
evaluate_model(args.model, args.test_data)
```

4. Kubeflow Pipeline Definition

```
1 # pipeline.py
2 from kfp import dsl
3 from kfp import components
4
5 def prepare data op():
6
       return dsl.ContainerOp(
7
           name='Prepare Data',
8
           image='python:3.8',
9
            command=['python', 'dataset_prep.py'],
10
            file outputs={
11
               'train': '/train.csv',
               'test': '/test.csv'
12
13
           }
14
       )
15
16
   def train_model_op(train_data):
17
       return dsl.ContainerOp(
18
            name='Train Model',
19
            image='python:3.8',
20
            command=['python', 'train.py'],
21
            arguments=['--data', train data],
22
            file_outputs={'model': '/model.pkl'}
23
24
25 def evaluate_model_op(model, test_data):
26
       return dsl.ContainerOp(
27
            name='Evaluate Model',
28
           image='python:3.8',
29
            command=['python', 'evaluate.py'],
30
            arguments=[
31
               '--model', model,
                '--test-data', test_data
32
33
34
            file_outputs={'metrics': '/metrics.json'}
35
36
37 @dsl.pipeline(
       name='Diabetes Prediction Pipeline',
38
39
       description='Train and evaluate a diabetes prediction model'
40 )
41 def diabetes_pipeline():
42
       data_prep = prepare_data_op()
43
       train = train_model_op(data_prep.outputs['train'])
44
       evaluate = evaluate_model_op(
45
           train.outputs['model'],
46
            data_prep.outputs['test']
47
48
49 # Compile pipeline
50 from kfp.compiler import Compiler
51 Compiler().compile(diabetes_pipeline, 'diabetes_pipeline.yaml')
```

5. Deploy Model to Azure ML

```
1 # deploy.py
2 from azureml.core import Workspace, Model, Environment
3 from azureml.core.webservice import AciWebservice
4 from azureml.core.model import InferenceConfig
5
6 def deploy_model(model_path):
7
        # Initialize workspace
8
       ws = Workspace.from_config()
9
10
        # Register model
       model = Model.register(
11
12
           workspace=ws,
13
           model path=model path,
            model name='diabetes-predictor'
14
15
       )
16
17
        # Create environment
18
       env = Environment.from_conda_specification(
19
            name='diabetes-env',
20
            file_path='environment.yml'
21
22
23
        # Define inference config
24
        inference config = InferenceConfig(
25
            entry_script='score.py',
            environment=env
26
27
28
29
        # Define deployment config
30
        deployment_config = AciWebservice.deploy_configuration(
31
            cpu_cores=1,
32
            memory gb=1,
33
            auth_enabled=True
34
        )
35
36
       # Deploy model
37
       service = Model.deploy(
38
           workspace=ws,
39
            name='diabetes-service',
40
            models=[model],
41
            inference_config=inference_config,
42
            deployment_config=deployment_config
43
44
45
        service.wait_for_deployment(show_output=True)
46
        return service.scoring_uri
47
48 if __name__ == '__main__':
        import argparse
49
50
       parser = argparse.ArgumentParser()
51
        parser.add argument('--model', type=str)
52
        args = parser.parse_args()
53
       deploy_model(args.model)
```

6. Scoring Script

```
1 # score.py
2 import json
```

```
3 import numpy as np
4 import pandas as pd
5 from azureml.core.model import Model
6 import pickle
7
8 def init():
9
     global model
10
     model_path = Model.get_model_path('diabetes-predictor')
11
       with open(model path, 'rb') as f:
           model = pickle.load(f)
12
13
14 def run(raw_data):
15
     try:
          data = json.loads(raw_data)
16
17
         data = pd.DataFrame(data)
18
         prediction = model.predict(data)
19
           return json.dumps({'prediction': prediction.tolist()})
20
       except Exception as e:
21
           return json.dumps({"error": str(e)})
```

Environment Configuration

Conda Environment

Usage Instructions

1. Set up infrastructure:

```
1 # Follow the setup steps above to create AKS cluster and deploy Kubeflow
```

2. Prepare environment:

```
1 # Create conda environment
2 conda env create -f environment.yml
3 conda activate diabetes-env
```

3. Run pipeline:

```
1 # Upload pipeline to Kubeflow
2 python pipeline.py
3
4 # Access Kubeflow UI
5 kubectl port-forward svc/istio-ingressgateway -n istio-system 8080:80
```

```
6 # Open browser at <a href="http://localhost:8080">http://localhost:8080>
```

4. Monitor deployment:

```
1 # Check pod status
2 kubectl get pods -n kubeflow
3
4 # Check service status
5 kubectl get services -n kubeflow
```

Monitoring

1. Azure Monitor Integration:

```
# Enable monitoring
from azureml.core import Workspace
from azureml.core.webservice import AciWebservice

ws = Workspace.from_config()
service = AciWebservice(ws, 'diabetes-service')

# Get logs
print(service.get_logs())

# Get metrics
print(service.get_metrics())
```

2. Kubeflow Dashboard:

- · Access metrics and logs through the Kubeflow UI
- Monitor pipeline runs and component status
- · Track model versions and deployments

This implementation provides a complete ML pipeline that:

- Uses the diabetes dataset as an example
- Implements data preparation, model training, and evaluation
- Deploys the model to Azure ML
- · Includes monitoring capabilities

The code is production-ready but may need adjustments based on your specific requirements and scale.