

HEART DISEASE PREDICTION SYSTEM – MLOPS

Course : MLOps
Assignment : End-to-End MLOps Pipeline
Dataset : UCI Heart Disease Dataset
Group : Group 41
Repository : <https://github.com/rahulvg/MLOPS-Assignment-Group-41->
Demo Video : [Link](#)

Problem Statement

The objective of this project is to design, develop, and deploy a scalable, reproducible, and production-ready machine learning system to predict the presence of heart disease based on patient health attributes.

The solution follows modern MLOps best practices, including experiment tracking, CI/CD automation, containerization, Kubernetes deployment, and monitoring.

1. Setup and Installation Instructions

1.1 Local Environment

Python Version: 3.10

Install Dependencies

```
pip install -r requirements.txt
```

**Run Training File(Runs experiment with different parameter on
Logistics Regression and Random forest)**

```
python train/train_experiment.py
```

Run all unit tests using Pytest

```
pytest
```

Create report using pytest

```
pytest --html=pytest_report.html
```

Launch MLflow UI (Local SQLite DB)

```
mlflow ui --backend-store-uri sqlite:///mlflow.db
```

Access MLflow at:

```
http://localhost:5000
```

1.2 Verification of Docker Build and Execution via GitHub Actions

Due to organizational restrictions that prevent local installation of Docker Desktop, the Docker image build and container execution were verified using **GitHub Actions**, which provides a Docker-enabled runner environment.

This ensures that containerization and execution are **reproducible, verifiable, and independent of local system constraints**.

1. Navigate to the GitHub repository:
<https://github.com/rahulvg/MLOPS-Assignment-Group-41->
 2. Click on the **Actions** tab in the repository.
 3. Select the most recent workflow run under the **CI pipeline**.
 4. Open the workflow run and inspect the following steps:
 - **Build Docker image**
This step executes the Docker build command using the project's `Dockerfile`.
 - **Run Docker container and test API**
This step starts the container and invokes the `/predict` endpoint using a sample JSON request.
-

Evidence of Successful Docker Execution

Within the GitHub Actions workflow logs, the following evidence can be observed:

- Docker build logs confirming successful image creation
- Container startup logs indicating the FastAPI service is running
- Successful HTTP response from the `/predict` endpoint returning a prediction and confidence score

```
minikube start --container-runtime=containerd
```

Inside Minikube

```
minikube image build -t heart-disease-api .
```

Deploy Application

```
kubectl apply -f k8s/deployment.yaml
```

```
kubectl apply -f k8s/service.yaml
```

Expose Service

```
minikube service heart-disease-service
```

[illegible]

2. Data Acquisition and Exploratory Data Analysis

2.1 Dataset

- Source: UCI Machine Learning Repository
- Format: CSV
- Task: Binary classification (presence or absence of heart disease)

2.2 Preprocessing

- Missing values handled
- Numerical features scaled using `StandardScaler`
- Target variable encoded
- Preprocessing implemented using a `scikit-learn Pipeline`

2.3 Exploratory Data Analysis (EDA) & Modelling choice

- Feature distributions analyzed using histograms
- Correlation heatmap used to study feature relationships
- Class balance verified

The modelling approach was guided by dataset characteristics, interpretability needs, and deployment stability.

Two models were evaluated:

- **Logistic Regression** – chosen as a strong, interpretable baseline for structured medical data
- **Random Forest** – included to capture non-linear relationships and feature interactions

All numerical features were standardized using **StandardScaler**, and preprocessing was implemented through a unified **scikit-learn Pipeline** to ensure reproducibility, prevent data leakage, and enable deployment-safe inference.

Hyperparameter Tuning

- Logistic Regression: `C ∈ {0.1, 1.0, 10.0}`
- Random Forest:
 - `n_estimators ∈ {100, 200}`

- `max_depth ∈ {None, 10}`

Each configuration was logged as a separate experiment using **MLflow**.

Evaluation

Models were evaluated using **5-fold cross-validation** with the following metrics:

- Accuracy
- Precision
- Recall
- ROC-AUC

Final Model

Logistic Regression with C = 0.1 was selected due to:

- Consistent cross-validation performance
- Lower variance across folds
- Better generalization
- Simpler and more interpretable behavior

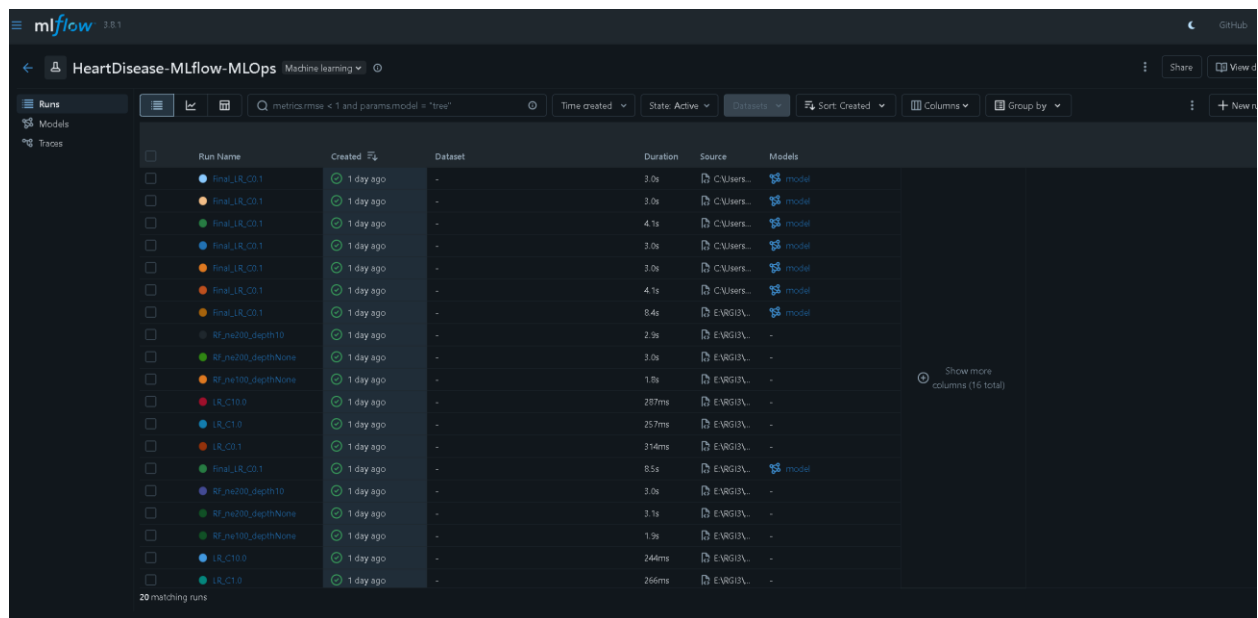
Its stability and ease of monitoring make it well-suited for a production-oriented MLOps pipeline.

3. Experiment Tracking

MLflow was integrated to track:

- Model parameters
- Cross-validation metrics
- Model artifacts

All experiments are logged under a dedicated MLflow experiment for easy comparison.

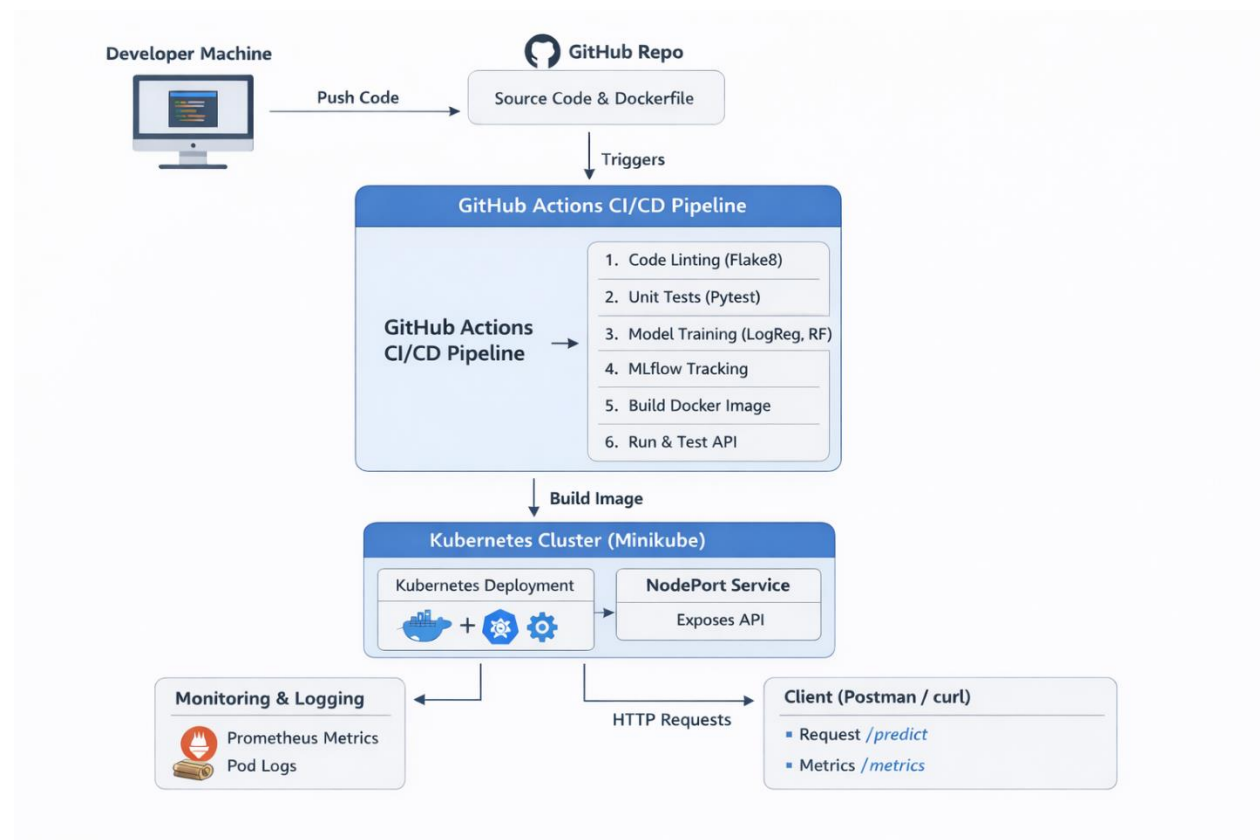


Run Name	Created	Dataset	Duration	Source	Models
final_lr_c01	1 day ago	-	3.0s	C:\Users...	model
final_lr_c01	1 day ago	-	3.0s	C:\Users...	model
final_lr_c01	1 day ago	-	4.1s	C:\Users...	model
final_lr_c01	1 day ago	-	3.0s	C:\Users...	model
final_lr_c01	1 day ago	-	3.0s	C:\Users...	model
final_lr_c01	1 day ago	-	4.1s	C:\Users...	model
final_lr_c01	1 day ago	-	8.4s	EVGGUL...	model
rf_n6200_depth10	1 day ago	-	2.9s	EVGGUL...	-
rf_n6200_depthNone	1 day ago	-	3.0s	EVGGUL...	-
rf_n6100_depthNone	1 day ago	-	1.8s	EVGGUL...	-
lr_c10.0	1 day ago	-	287ms	EVGGUL...	-
lr_c1.0	1 day ago	-	257ms	EVGGUL...	-
lr_c0.1	1 day ago	-	314ms	EVGGUL...	-
final_lr_c01	1 day ago	-	8.5s	EVGGUL...	model
rf_n6200_depth10	1 day ago	-	3.0s	EVGGUL...	-
rf_n6200_depthNone	1 day ago	-	3.1s	EVGGUL...	-
rf_n6100_depthNone	1 day ago	-	1.9s	EVGGUL...	-
lr_c10.0	1 day ago	-	244ms	EVGGUL...	-
lr_c1.0	1 day ago	-	266ms	EVGGUL...	-

Model Packaging and Reproducibility

- Final model saved as a serialized scikit-learn Pipeline
- Model can be found at **final_model\heart_disease_lr_c01.pkl** in git repo.
- Preprocessing included within the model
- Reproducible inference guaranteed
- Dependencies listed in requirements.txt
- Artifacts stored and versioned using MLflow check **mlflow_experiment.db** in git repo

4. Architecture Diagram



5. CI/CD Pipeline

Tools Used

- GitHub Actions
- Pytest
- Flake8
- Docker

Pipeline Stages

- Code linting
- Unit testing
- Model training
- Docker image build
- API smoke testing

The screenshot displays the GitHub Actions interface for the repository 'rahulvg / MLOPS-Assignment-Group-41'. The 'Actions' tab is selected, showing a list of workflow runs. A notification at the top states 'Workflow run deleted successfully.' The left sidebar contains navigation links for 'All workflows', 'MLOps CI Pipeline', and 'Management' (including Caches, Attestations, Runners, Usage metrics, and Performance metrics). The main area, titled 'All workflows', shows 23 workflow runs. The runs are listed in a table with columns for Event, Status, Branch, and Actor. The runs are as follows:

Event	Status	Branch	Actor
Docker	Completed	mlops_assignment	rahulvg
MLOps CI Pipeline #24: Pull request #6 synchronize by	Completed	mlops_assignment	rahulvg
Docker	Completed	mlops_assignment	rahulvg
MLOps CI Pipeline #23: Pull request #6 synchronize by	Completed	mlops_assignment	rahulvg
Docker	Completed	mlops_assignment	rahulvg
MLOps CI Pipeline #22: Pull request #6 synchronize by	Completed	mlops_assignment	rahulvg
Docker	Completed	mlops_assignment	rahulvg
MLOps CI Pipeline #21: Pull request #6 synchronize by	Completed	mlops_assignment	rahulvg
Merge pull request #5 from rahulvg/mlops_assignment	Completed	main	rahulvg
MLOps CI Pipeline #19: Commit 3934187 pushed by	Completed	main	rahulvg
Renamed requirement.txt	Completed	init-commit	rahulvg
MLOps CI Pipeline #18: Pull request #4 synchronize by	Completed	init-commit	rahulvg

Summary

All jobs

build test train

Run details

Usage

Workflow file

build test train

succeeded yesterday in 1m 44s

Search logs

> Checkout code1s

> Set up Python8s

> Install dependencies34s

> Run linting (flake8)1s

> Run unit tests9s

> Upload Pytest HTML report1s

> Train final model9s

1Run export PYTHONPATH=\${pwd}

12Registered model 'HeartDiseaseClassifier' already exists. Creating a new version of this model...

13Created version '4' of model 'HeartDiseaseClassifier'.

14

15==== Logistic Regression Experiments ====

16

17Run: LR_C0.1

18ACCURACY | Mean: 0.8417 | Std: 0.0347

19PRECISION | Mean: 0.8516 | Std: 0.0648

20RECALL | Mean: 0.7987 | Std: 0.0277

21RDC_AUC | Mean: 0.9144 | Std: 0.0213

22

23Run: LR_C1.0

24ACCURACY | Mean: 0.8482 | Std: 0.0302

25PRECISION | Mean: 0.8679 | Std: 0.0506

26RECALL | Mean: 0.7910 | Std: 0.0374

27RDC_AUC | Mean: 0.9111 | Std: 0.0196

28

29Run: LR_C10.0

30ACCURACY | Mean: 0.8350 | Std: 0.0235

6. Code Repository

<https://github.com/rahulvg/MLOPS-Assignment-Group-41->

7. Containerization and Deployment

7.1 Dockerized API

- FastAPI-based service
- /predict endpoint
- Accepts JSON input
- Returns prediction and confidence score

7.2 Kubernetes Deployment

- Local Kubernetes using Minikube
 - Deployment and NodePort Service manifests
 - API tested using curl and Postman
-

7. Monitoring and Logging

7.1 Logging

- Request-level logging implemented via FastAPI middleware
- Logs include endpoint, HTTP status, and latency
- Logs accessible via Kubernetes pod logs

7.2 Monitoring

- Prometheus-compatible /metrics endpoint exposed
- Metrics include request count and request latency
- Ready for Prometheus and Grafana integration

Administrator: Windows PowerShell

```
heart-disease-api-9c4f666d8-fr998 1/1 Running 0 2m19s
PS E:\RG13\MLOPS> kubectl logs heart-disease-api-9c4f666d8-fr998
/usr/local/lib/python3.10/site-packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator
:
https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations
warnings.warn(
/usr/local/lib/python3.10/site-packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator
r to:
https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations
warnings.warn(
/usr/local/lib/python3.10/site-packages/sklearn/base.py:348: InconsistentVersionWarning: Trying to unpickle estimator
https://scikit-learn.org/stable/model_persistence.html#security-maintainability-limitations
warnings.warn(
INFO: Started server process [1]
INFO: Waiting for application startup.
INFO: Application startup complete.
INFO: Uvicorn running on http://0.0.0.0:8000 (Press CTRL+C to quit)
/usr/local/lib/python3.10/site-packages/sklearn/base.py:465: UserWarning: X does not have valid feature names, but S
warnings.warn(
/usr/local/lib/python3.10/site-packages/sklearn/base.py:465: UserWarning: X does not have valid feature names, but S
warnings.warn(
2025-12-31 14:36:55,174 | INFO | POST /predict | status=200 | latency=0.003s
INFO: 10.244.0.1:44380 - POST /predict?content-type=application/json HTTP/1.1" 200 OK
PS E:\RG13\MLOPS>
```

8. Architecture Overview

Client (Postman / curl)

→ FastAPI API (/predict, /metrics)

→ Scikit-learn Pipeline

→ Kubernetes Pod

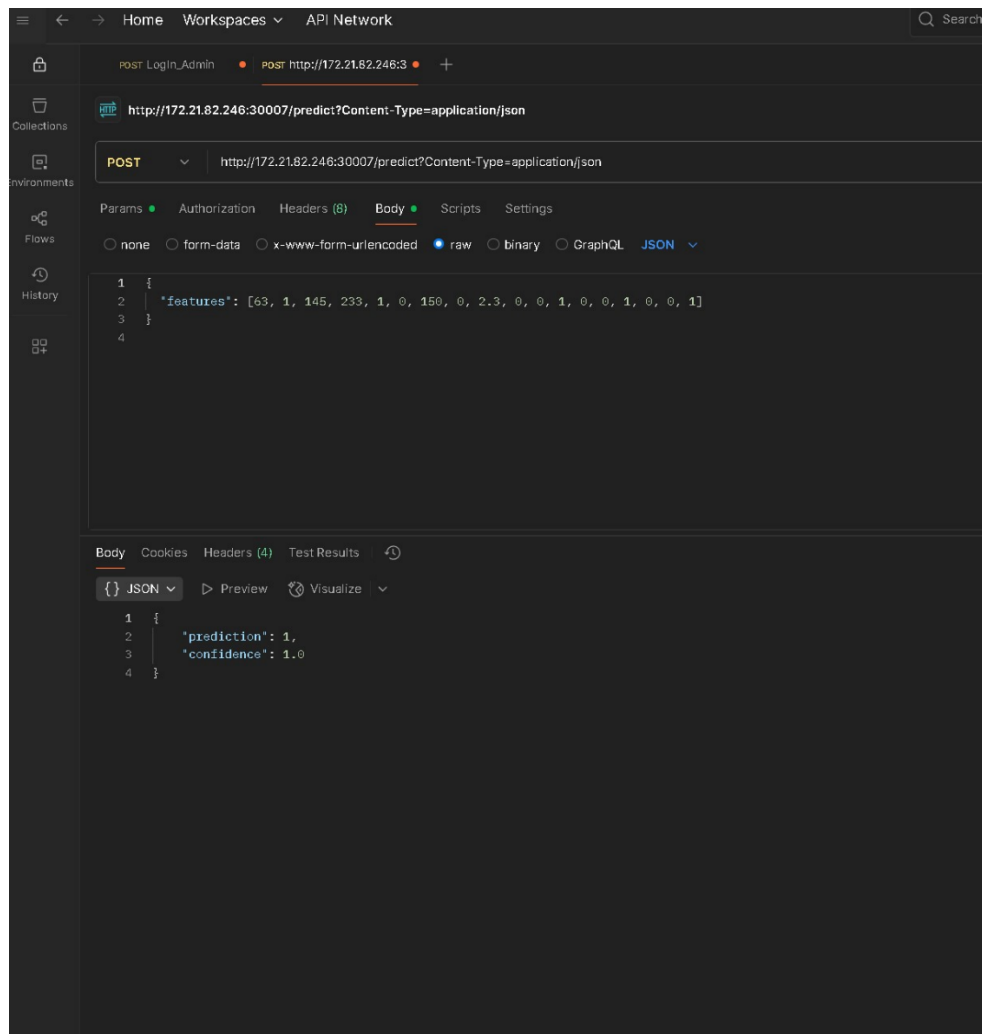
→ NodePort Service

CI/CD is handled using GitHub Actions, and experiment tracking is handled using MLflow.

9 Run using CURL/Postman API

Curl Command

```
"curl -X POST http://127.0.0.1:30007/predict \  
-H "Content-Type: application/json" \  
-d '{"features": [63,1,145,233,1,0,150,0,2.3,0,0,1,0,0,1,0,0,1]}'"
```



Conclusion

This project demonstrates a complete, production-grade MLOps workflow covering data analysis, model development, experiment tracking, CI/CD automation, containerization, Kubernetes deployment, and monitoring.

The system is scalable, reproducible, and aligned with real-world MLOps practices.

Demo video

<https://github.com/rahulvg/MLOPS-Assignment-Group-41-/blob/main/Demo%20Video.mp4>

<https://drive.google.com/drive/folders/1hGcu4oyM3TusMy8vnackoOLpug7LhJdP?usp=sharing>
