

Smart Health Analysis

A machine learning project that predicts potential health conditions using common blood biomarkers through a simple Flask web app.

What It Does

- Lets users input health markers through a web form.
 - Uses trained machine learning models to classify potential conditions.
 - Supports multiple health categories: **Fit**, **Anemia**, **Hypertension**, **Diabetes**, and **High Cholesterol**.
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Biomarkers Used

- HbA1C
 - Systolic Blood Pressure
 - Diastolic Blood Pressure
 - LDL Cholesterol
 - HDL Cholesterol
 - Triglycerides
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Project Files

```
Smart_Health_Analysis/  
├─ app.py                # Main Flask application  
├─ for_deploy.py        # Alternative entrypoint for deployment  
├─ templates/           # Web app HTML templates  
├─ health_markers_dataset.csv # Training dataset  
├─ health-markers-dataset.ipynb # Data exploration and model training notebook  
├─ Voting_health_model.pkl # Trained ensemble model  
├─ health_markers_dataset_model.pkl # Baseline model  
└─ .venv/               # Virtual environment (local only)
```

Getting Started

1. Clone the Repository

```
git clone https://github.com/MohammedSameh2/Smart_Health_Analysis.git
cd Smart_Health_Analysis
```

2. Create and Activate a Virtual Environment

Windows

```
python -m venv .venv
.venv\Scripts\activate
```

macOS/Linux

```
python3 -m venv .venv
source .venv/bin/activate
```

3. Install Dependencies

If `requirements.txt` is available:

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

Or install manually:

```
pip install flask scikit-learn pandas numpy joblib matplotlib
```

4. Run the Application

```
python app.py
```

Open your browser at **`http://127.0.0.1:5000/`**.

The Models

- **Baseline model** (`health_markers_dataset_model.pkl`) trained directly on the dataset.

- **Ensemble model** (`Voting_health_model.pkl`) that combines multiple algorithms for better accuracy.
 - Training workflow and experiments are detailed in the Jupyter notebook.
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Example Usage

```
import joblib
import numpy as np

model = joblib.load('Voting_health_model.pkl')
example = np.array([[6.2, 130, 82, 160, 45, 180]])
print(model.predict(example))
```

Credits

Developed by **Mohamed Sameh** using Python, Flask, and scikit-learn.