

## Main code Of binary text based model loading

```
from fastapi import FastAPI
from fastapi.middleware.cors import CORSMiddleware
from pydantic import BaseModel
import tensorflow as tf
import numpy as np
# Initialize FastAPI app
app = FastAPI()
# Enable CORS to allow requests from Flutter
app.add_middleware(
    CORSMiddleware,
    allow_origins=["*"], # Allow all origins for testing, restrict in production
    allow_credentials=True,
    allow_methods=["*"], # Allow all HTTP methods
    allow_headers=["*"], # Allow all headers
)
# Load the model
try:
    model = tf.keras.models.load_model("heart_disease_model.h5")
    print("Model loaded successfully!")
except Exception as e:
    print(f"Error loading model: {e}")
# Input data model for request
class InputData(BaseModel):
    age: int
    sex: int
    cp: int
    trestbps: int
    chol: int
    fbs: int
    restecg: int
    thalach: int
    exang: int
    oldpeak: float
    slope: int
    ca: int
    thal: int
# Define a route to check the server status
@app.get("/")
def read_root():
    return {"message": "Welcome to the Heart Disease Predictor!"}
# Prediction route
@app.post("/predict")
def predict(data: InputData):
    try:
        # Prepare input data for the model
        input_data = np.array([
            data.age, data.sex, data.cp, data.trestbps, data.chol, data.fbs,
            data.restecg, data.thalach, data.exang, data.oldpeak, data.slope,
            data.ca, data.thal
        ])
        # Make the prediction
        prediction = model.predict(input_data)
        result = "Heart Disease Detected" if prediction[0][0] > 0.5 else "No
Heart Disease Detected"

        return {"prediction": result}
    except Exception as e:
        return {"error": str(e)}
```

## If String type value available handle like that

```
from fastapi import FastAPI
from pydantic import BaseModel
from fastapi.middleware.cors import CORSMiddleware
import tensorflow as tf
import numpy as np

# Initialize FastAPI app
app = FastAPI()

# Enable CORS to allow requests from Flutter
app.add_middleware(
    CORSMiddleware,
    allow_origins=["*"], # Allow all origins for testing, restrict in production
    allow_credentials=True,
    allow_methods=["*"], # Allow all HTTP methods
    allow_headers=["*"], # Allow all headers
)

# Load your model (change the path as needed)
try:
    model = tf.keras.models.load_model("your_model.h5") # Replace with your model file
    print("Model loaded successfully!")
except Exception as e:
    print(f"Error loading model: {e}")

# Input data model for request
class InputData(BaseModel):
    age: int
    bp: int # Blood pressure
    sg: float # Specific gravity
    al: int # Albumin
    su: int # Sugar
    rbc: str # Red blood cells (categorical)
    pc: str # Pus cell (categorical)
    pcc: str # Pus cell clumps (categorical)
    ba: str # Bacteria (categorical)
    bgr: int # Blood glucose random
    bu: float # Blood urea
    sc: float # Serum creatinine
    sod: float # Sodium
    pot: float # Potassium
    hemo: float # Hemoglobin
    pcv: float # Packed cell volume
    wc: float # White blood cell count
    rc: float # Red cell count
    htn: str # Hypertension (categorical)
    dm: str # Diabetes mellitus (categorical)
    cad: str # Coronary artery disease (categorical)
    appet: str # Appetite (categorical)
    pe: str # Pedal edema (categorical)
    ane: str # Anemia (categorical)

# Define a route to check the server status
@app.get("/")
def read_root():
    return {"message": "Welcome to the CKD Predictor!"}

# Prediction route
@app.post("/predict")
def predict(data: InputData):
```

```

try:
    # Prepare input data for the model
    # Convert categorical features to numeric (e.g., one-hot encoding or label encoding)
    # Example encoding (replace these with the actual encoding used in your model)
    rbc_encoded = 1 if data.rbc == "present" else 0
    pc_encoded = 1 if data.pc == "present" else 0
    pcc_encoded = 1 if data.pcc == "present" else 0
    ba_encoded = 1 if data.ba == "present" else 0
    htn_encoded = 1 if data.htn == "yes" else 0
    dm_encoded = 1 if data.dm == "yes" else 0
    cad_encoded = 1 if data.cad == "yes" else 0
    appet_encoded = 1 if data.appet == "good" else 0
    pe_encoded = 1 if data.pe == "yes" else 0
    ane_encoded = 1 if data.ane == "yes" else 0

    input_data = np.array([[
        data.age,
        data.bp,
        data.sg,
        data.al,
        data.su,
        rbc_encoded,
        pc_encoded,
        pcc_encoded,
        ba_encoded,
        data.bgr,
        data.bu,
        data.sc,
        data.sod,
        data.pot,
        data.hemo,
        data.pcv,
        data.wc,
        data.rc,
        htn_encoded,
        dm_encoded,
        cad_encoded,
        appet_encoded,
        pe_encoded,
        ane_encoded,
    ]])

    # Make the prediction
    prediction = model.predict(input_data)

    # Customize this logic based on your model's output
    result = "CKD Detected" if prediction[0][0] > 0.5 else "No CKD Detected"

    return {"prediction": result}
except Exception as e:
    return {"error": str(e)}

```

## Environment Setup

```

# Create a virtual environment
python -m venv heart-disease-env

```

```

# Activate the virtual environment

```

```

# Windows
heart-disease-env\Scripts\activate

```

```
# macOS/Linux  
sourceheart-disease-env/bin/activate
```

```
# Install required packages  
pip install fastapi uvicorn tensorflow numpy
```