## Water portability checker

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Model_train.py:
import pandas as pd
import xgboost as xgb
from sklearn.model selection import train test split
from sklearn.preprocessing import StandardScaler
from sklearn.metrics import accuracy_score
from imblearn.over_sampling import SMOTE
import pickle
df = pd.read_csv('dataset (1).csv')
df = df.fillna(df.mean())
X = df.drop('Potability', axis=1)
y = df['Potability']
scaler = StandardScaler()
X_scaled = scaler.fit_transform(X)
X_train, X_test, y_train, y_test = train_test_split(
  X_scaled, y, test_size=0.2, stratify=y, random_state=42
smote = SMOTE(random_state=42)
X_train, y_train = smote.fit_resample(X_train, y_train)
```

```
model = xgb.XGBClassifier(
  n_estimators=500,
  max depth=10,
  learning_rate=0.01,
  subsample=0.8,
  colsample_bytree=0.8,
  use_label_encoder=False,
  eval_metric='logloss'
)
model.fit(X_train, y_train)
preds = model.predict(X_test)
acc = accuracy_score(y_test, preds)
print(f'Accuracy: {acc}')
with open('water_model.pkl', 'wb') as f:
  pickle.dump(model, f)
with open('scaler.pkl', 'wb') as f:
  pickle.dump(scaler, f)
Main.py:
from fastapi import FastAPI
from pydantic import BaseModel
```

```
import pickle
import numpy as np
with open('water model.pkl', 'rb') as f:
  model = pickle.load(f)
with open('scaler.pkl', 'rb') as f:
  scaler = pickle.load(f)
app = FastAPI()
class WaterInput(BaseModel):
  ph: float
  Hardness: float
  Solids: float
  Chloramines: float
  Sulfate: float
  Conductivity: float
  Organic_carbon: float
  Trihalomethanes: float
  Turbidity: float
@app.post("/predict")
def predict(data: WaterInput):
  input_data = np.array([[
    data.ph, data.Hardness, data.Solids, data.Chloramines,
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data.Sulfate, data.Conductivity, data.Organic_carbon,
    data.Trihalomethanes, data.Turbidity
 ]])
  input scaled = scaler.transform(input data)
  pred = model.predict(input_scaled)[0]
  return {"Potable": bool(pred)}
App.py
import streamlit as st
import pandas as pd
import numpy as np
import pickle
import plotly.express as px
st.set_page_config(
  page_title="Water Quality Analysis",
  page_icon=" \( \bigcup \)",
 layout="wide"
)
st.markdown("""
<style>
  .main {
    padding: 20px;
  }
  .stButton>button {
    width: 100%;
```

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background-color: #4CAF50;
  color: white;
  height: 3em;
  margin-top: 20px;
  border-radius: 10px;
  font-size: 18px;
  font-weight: bold;
}
.stButton>button:hover {
  background-color: #45a049;
}
.stSlider {
  padding-top: 10px;
  padding-bottom: 10px;
}
.title {
  font-size: 48px;
  font-weight: bold;
  color: #1e88e5;
  text-align: center;
  margin-bottom: 30px;
  text-shadow: 2px 2px 4px rgba(0,0,0,0.1);
}
.subtitle {
  font-size: 24px;
  color: #424242;
```

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margin-bottom: 20px;
    text-align: center;
  }
  .result-box {
    padding: 20px;
    border-radius: 10px;
    margin: 20px 0;
    text-align: center;
    font-size: 24px;
    font-weight: bold;
  }
  .info-box {
    background-color: #f8f9fa;
    padding: 20px;
    border-radius: 10px;
    margin-bottom: 20px;
  }
</style>
""", unsafe_allow_html=True)
@st.cache_resource
def load_model():
  with open('water_model.pkl', 'rb') as f:
    model = pickle.load(f)
  with open('scaler.pkl', 'rb') as f:
    scaler = pickle.load(f)
```

```
return model, scaler
```

```
st.markdown("<h1 class='title'>  Water Potability Analysis</h1>", unsafe_allow_html=True)
st.markdown("""
<div class='info-box'>
  Analyze your water quality parameters to determine if the water is safe for drinking.
    This Al-powered tool uses advanced machine learning to assess water potability based on
various chemical and physical properties.
  </div>
""", unsafe allow html=True)
try:
  model, scaler = load_model()
  col1, col2, col3 = st.columns(3)
  with col1:
    st.markdown("<h3 style='text-align: center; color: #1e88e5;'>Chemical Properties</h3>",
unsafe_allow_html=True)
    ph = st.slider('pH Level', min_value=0.0, max_value=14.0, value=7.0)
    sulfate = st.slider('Sulfate', min_value=0.0, max_value=500.0, value=250.0)
    chloramines = st.slider('Chloramines', min value=0.0, max value=15.0, value=7.0)
  with col2:
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st.markdown("<h3 style='text-align: center; color: #1e88e5;'>Physical Properties</h3>",
unsafe allow html=True)
    hardness = st.slider('Hardness', min value=0.0, max value=500.0, value=200.0)
    solids = st.slider('Total Dissolved Solids', min_value=0.0, max_value=1000.0, value=500.0)
    conductivity = st.slider('Conductivity', min_value=0.0, max_value=1000.0, value=500.0)
  with col3:
    st.markdown("<h3 style='text-align: center; color: #1e88e5;'>Organic Properties</h3>",
unsafe allow html=True)
    organic_carbon = st.slider('Organic Carbon', min_value=0.0, max_value=30.0, value=15.0)
    trihalomethanes = st.slider('Trihalomethanes', min value=0.0, max value=150.0,
value=75.0)
    turbidity = st.slider('Turbidity', min value=0.0, max value=10.0, value=5.0)
  input_data = pd.DataFrame({
    'ph': [ph],
    'Hardness': [hardness],
    'Solids': [solids],
    'Chloramines': [chloramines],
    'Sulfate': [sulfate],
    'Conductivity': [conductivity],
    'Organic carbon': [organic carbon],
    'Trihalomethanes': [trihalomethanes],
    'Turbidity': [turbidity]
  })
  if st.button('Analyze Water Quality'):
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with st.spinner('Analyzing water quality...'):
  input scaled = scaler.transform(input data)
  prediction = model.predict(input scaled)
  probability = model.predict proba(input scaled)
  st.markdown("<br>", unsafe allow html=True)
  result col, chart col = st.columns(2)
  with result col:
    if prediction[0] == 1:
      st.markdown(f"""
        <div class='result-box' style='background-color: #e8f5e9; color: #2e7d32;'>
           POTABLE WATER<br>
          Confidence: {probability[0][1]:.1%}
        </div>
      """, unsafe allow html=True)
    else:
      st.markdown(f"""
        <div class='result-box' style='background-color: #ffebee; color: #c62828;'>
           ♠ NON-POTABLE WATER<br>
          Confidence: {probability[0][0]:.1%}
        </div>
      """, unsafe_allow_html=True)
  with chart col:
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names=['Non-Potable', 'Potable'],
               title='Prediction Confidence',
               hole=0.6,
               color_discrete_sequence=['#ef5350', '#66bb6a'])
         fig.update_layout(
           title_x=0.5,
           title font size=20,
           showlegend=True,
           legend=dict(orientation="h", yanchor="bottom", y=1.02, xanchor="center", x=0.5)
        )
         st.plotly_chart(fig)
      st.markdown("<h3 style='text-align: center; color: #1e88e5;'>Water Quality
Parameters</h3>", unsafe_allow_html=True)
      param cols = st.columns(3)
      param ranges = {
         'pH': {'min': 0, 'max': 14, 'optimal min': 6.5, 'optimal max': 8.5},
         'Hardness': {'min': 0, 'max': 500, 'optimal_min': 50, 'optimal_max': 300},
         'TDS': {'min': 0, 'max': 1000, 'optimal min': 100, 'optimal max': 500}
      }
      with param_cols[0]:
         fig = px.bar([{'Parameter': 'pH', 'Value': ph}],
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fig = px.pie(values=[probability[0][0], probability[0][1]],

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x='Parameter', y='Value',
               title='pH Level',
               color discrete sequence=['#1e88e5'])
        fig.add hline(y=6.5, line dash="dash", line color="green", annotation text="Min
Safe")
        fig.add hline(y=8.5, line dash="dash", line color="red", annotation text="Max Safe")
        fig.update layout(title x=0.5)
        st.plotly_chart(fig)
      with param_cols[1]:
        fig = px.bar([{'Parameter': 'TDS', 'Value': solids}],
               x='Parameter', y='Value',
               title='Total Dissolved Solids',
               color discrete sequence=['#1e88e5'])
        fig.add_hline(y=500, line_dash="dash", line_color="red", annotation_text="Max Safe")
        fig.update layout(title x=0.5)
        st.plotly chart(fig)
      with param cols[2]:
        fig = px.bar([{'Parameter': 'Hardness', 'Value': hardness}],
               x='Parameter', y='Value',
               title='Water Hardness',
               color_discrete_sequence=['#1e88e5'])
        fig.add hline(y=300, line dash="dash", line color="red", annotation text="Max Safe")
        fig.update_layout(title_x=0.5)
        st.plotly chart(fig)
```

## except Exception as e:

st.error(f"An error occurred: {str(e)}")

st.info("Please make sure the model files (water\_model.pkl and scaler.pkl) are present in the same directory.")