

# train-models

December 31, 2025

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[2]: # Importing our libraries
import pandas as pd
import numpy as np
import pickle
import os
from sklearn.model_selection import train_test_split
from sklearn.multioutput import MultiOutputClassifier
from sklearn.ensemble import RandomForestClassifier
from sklearn.svm import SVC
from sklearn.neighbors import KNeighborsClassifier
import xgboost as xgb

[3]: # 1. SETUP
os.makedirs("models", exist_ok=True)

[4]: # 2. LOAD DATA (Prioritize the Cleaned "No Sleep" File)
if os.path.exists("final_dataset_no_sleep.csv"):
    print(" Loading pre-cleaned dataset: 'final_dataset_no_sleep.csv'")
    df = pd.read_csv("final_dataset_no_sleep.csv")

    # Since the file is already cleaned, we just take all Q columns
    feature_cols = [c for c in df.columns if "Q" in c]

else:
    # Fallback: Load original/corrected and clean in-memory
    filename = "corrected_mental_health_dataset.csv"
    if not os.path.exists(filename):
        filename = "synthetic_college_mental_health_dataset_v3_with_severity.
        ↪csv"
        print(f" Clean file not found. Using '{filename}' and cleaning_
        ↪in-memory.")
    else:
        print(f" Clean file not found. Using '{filename}' and cleaning_
        ↪in-memory.")

    df = pd.read_csv(filename)
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# A. Apply Clinical Hierarchy (Safety Check)
overlap_mask = (df['ASD_risk'] == 1) & (df['SPCD_risk'] == 1)
df.loc[overlap_mask, 'SPCD_risk'] = 0

# B. REMOVE SLEEP QUESTIONS (Madam's Requirement)
all_q_cols = [c for c in df.columns if "Q" in c]
# Filter out any column containing "CONF"
feature_cols = [c for c in all_q_cols if "CONF" not in c]

print(f"Training on {len(feature_cols)} features (27 Expected).")

risk_cols = ["ADHD_risk", "ASD_risk", "SPCD_risk", "DEP_risk", "ANX_risk"]

X = df[feature_cols]
y = df[risk_cols]

```

Loading pre-cleaned dataset: 'final\_dataset\_no\_sleep.csv'  
Training on 27 features (27 Expected).

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[5]: # 3. SPLIT DATA
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.2,
    ↪random_state=42)

# Save Test Data for the Report Generator
pickle.dump((X_test, y_test), open("models/test_data.pkl", "wb"))
print("Saved 'models/test_data.pkl' for graph generation.")

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Saved 'models/test\_data.pkl' for graph generation.

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[6]: # 4. TRAIN ALL 4 RISK MODELS
models = {
    "RF": MultiOutputClassifier(RandomForestClassifier(n_estimators=100,
    ↪random_state=42)),
    # Removed use_label_encoder=False to fix UserWarning
    "XGB": MultiOutputClassifier(xgb.XGBClassifier(eval_metric='logloss',
    ↪random_state=42)),
    "SVM": MultiOutputClassifier(SVC(kernel='linear', probability=True,
    ↪random_state=42)),
    "KNN": MultiOutputClassifier(KNeighborsClassifier(n_neighbors=5))
}

for name, model in models.items():
    print(f"Training {name} Risk Model...")
    model.fit(X_train, y_train)
    pickle.dump(model, open(f"models/{name}_risk.pkl", "wb"))

```

Training RF Risk Model...

Training XGB Risk Model...  
Training SVM Risk Model...  
Training KNN Risk Model...

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[7]: # 5. TRAIN SEVERITY MODELS (Random Forest)
SEVERITY_MAPPING = {'Low': 0, 'Medium': 1, 'High': 2}
severity_map_cols = {
    "ADHD": "ADHD_severity", "ASD": "ASD_severity", "SPCD": "SPCD_severity",
    "DEP": "DEP_severity", "ANX": "ANX_severity"
}

print("\nTraining Severity Models (RF)...")
for disorder, sev_col in severity_map_cols.items():
    risk_col = f"{disorder}_risk"

    # Train only on rows where the disorder exists
    sev_df = df[df[risk_col] == 1].copy()

    X_sev = sev_df[feature_cols] # 27 cols
    y_sev = sev_df[sev_col].map(SEVERITY_MAPPING).fillna(0)

    rf_sev = RandomForestClassifier(n_estimators=100, max_depth=8,
    random_state=42)
    rf_sev.fit(X_sev, y_sev)
    pickle.dump(rf_sev, open(f"models/rf_{disorder}_sev.pkl", "wb"))

print("\n All models retrained successfully!")
```

Training Severity Models (RF)...

All models retrained successfully!