

Unsupervised Detection of Anomalous Wellness Profiles: Identifying Statistical Outliers in Self-Reported Mental Health and Lifestyle Data

Collaborative Final Project

CSST101 – Machine Learning

CSST102 – Knowledge Representation and Reasoning

Submitted by: Group Name: Group 2

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PROJECT OVERVIEW

Combine an Isolation Forest anomaly detector with rule-based reasoning to identify individuals with unusual wellness indicators and produce risk-level recommendations and next-step actions. Objectives

OBJECTIVES

General Objective: Detect and prioritize potential mental-health risk cases using a hybrid ML + knowledge-based system.

Specific Objectives:

- Build an unsupervised anomaly detection model to score wellness-related records.
- Define knowledge-based rules to translate anomaly signals into human-understandable risk levels.
- Produce an interpretable report listing top anomalies and recommended actions.
- Package model and pipeline for simple deployment and repeatable evaluation.

SYSTEM ARCHITECTURE

User Input → Preprocessing → Isolation Forest (ML) → KRR Rules (CSST102) → Final Risk Level → Recommendations

Report Artifacts: Model saved as output/isolation_forest.joblib, anomaly outputs in output/anomaly_scores.csv and output/top_anomalies.csv **Machine Learning Component (CSST101)**

MACHINE LEARNING COMPONENT (CSST101)

- **Algorithm Used:** Isolation Forest (unsupervised anomaly detection)
- **Dataset:** See archive (1)/Mental Health Dataset.csv/Mental%20Health%20Dataset.csv for raw records
- **Model Evaluation:** Use AUC/precision@k where labeled data available, plus inspection of top anomalies and cluster summaries in output/

MACHINE LEARNING PIPELINE

- **Data Collection:** Survey/CSV dataset containing wellness indicators and demographic fields (source: archive (1)/Mental Health Dataset.csv).
- **Data Preprocessing:** Handle missing values, normalize numeric features, encode categorical features, remove low-variance columns, engineer aggregated wellness scores.
- **Model Training:** Train IsolationForest on cleaned features; tune `n_estimators`, `contamination`, and `max_samples` via cross-validation or domain-guided selection.
- **Model Evaluation:** Validate using known anomalies (if any), inspect distribution of anomaly scores, check stability across folds, review output/cluster_summary.md.
- **Model Deployment:** Save model to output/isolation_forest.joblib; provide simple inference API or notebook cell in anomaly_analysis.ipynb.

DATASET DESCRIPTION

- **Dataset Type:** Tabular CSV (survey/clinical indicators)
- **Number of Records:** See archive (1)/Mental Health Dataset.csv/Mental%20Health%20Dataset.csv for the exact count
- **Target Variable:** No explicit label (unsupervised); target concept = anomalous/unusual wellness profile

KNOWLEDGE REPRESENTATION & REASONING (CSST102)

- **Rule 1:** IF `anomaly_score` ≥ 0.85 THEN Risk = High AND Recommend = "Clinical follow-up within 48 hours"
- **Rule 2:** IF $0.6 \leq \text{anomaly_score} < 0.85$ AND `severe_symptom_flag` = True THEN Risk = High AND Recommend = "Contact support hotline"
- **Rule 3:** IF $0.4 \leq \text{anomaly_score} < 0.6$ THEN Risk = Medium AND Recommend = "Schedule clinician review"
- **Rule 4:** IF `anomaly_score` < 0.4 AND `no_concerning_flags` THEN Risk = Low AND Recommend = "Routine monitoring"
- **Rule 5:** IF `demographic_vulnerability` = True AND `anomaly_score` ≥ 0.6 THEN Elevate Risk by one level AND Recommend = "Prioritized outreach"

HYBRID DECISION LOGIC

Logic: Compute anomaly score from Isolation Forest (0–1), apply deterministic KRR thresholds above, and combine with domain flags (e.g., `severe_symptom_flag`, `demographic_vulnerability`) using simple weighted rules to produce final Risk Level (Low/Medium/High) and corresponding recommendation text.

Aggregation: If multiple rules trigger, pick the highest risk; include all matching rule explanations in the report.

SYSTEM FEATURES

- Wellness risk prediction: Detects anomalous profiles.
- Rule-based recommendations: Maps scores to actionable steps.
- Reports & CSV outputs: `output/anomaly_report.md`, `output/top_anomalies.csv`.
- Model persistence: `output/isolation_forest.joblib`.
- Notebook & API-ready: `anomaly_analysis.ipynb` for interactive use.

TESTING AND EVALUATION

Test Case 1 | Input Summary: Synthetic record with extreme scores | Expected Output: $\text{anomaly_score} \geq 0.85 \rightarrow \text{Risk} = \text{High}$

Test Case 2 | Input Summary: Mild deviations, no flags | Expected Output: $\text{anomaly_score} \sim 0.5 \rightarrow \text{Risk} = \text{Medium}$

Test Case 3 | Input Summary: Normal profile | Expected Output: $\text{anomaly_score} < 0.4 \rightarrow \text{Risk} = \text{Low}$
Validation: Compare top anomalies with domain expert review; run `tests/test_pipeline.py` to verify end-to-end pipeline.

CONCLUSION

Outcome: A lightweight hybrid system that flags unusual wellness records, provides interpretable risk levels, and outputs recommendations for follow-up.

GROUP CONTRIBUTION

Member Name | Contribution

Dela Paz, Zyril - Main coder, Proof Reader

Delos Reyes, Axcel - Debugger, Lead Documenter

Hornilla, John Benedict - Dataset provider, Lead Presenter

REFERENCES

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