

Patient Readmission Prediction using H2O AutoML

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Introduction

Problem Statement:

• Hospital readmission prediction is critical for resource allocation, patient care, and cost reduction.

Objective:

Use H2O AutoML to automate model selection and improve prediction accuracy.

Key Result:

Best-performing model (GBM) achieved 99% accuracy, AUC = 1.0.

Literature Review



Traditional Approaches:

Logistic Regression, Decision
Trees – struggled with complex medical data.

Advancements:

ML models like XGBoost,
Random Forests, GBM improve predictions.

Why AutoML?

 Automates model selection & tuning, making healthcare predictions more accessible.

Dataset & Preprocessing

Dataset: elCU Collaborative Research Database (PhysioNet).

Key Tables Used: Patient, AdmissionDX, Diagnosis, Lab, Treatment.

Preprocessing Steps:

- Removed duplicates, optimized data types, merged tables.
- Dropped medication table due to high memory usage.
- Final dataset: 1000 rows, 14 columns.

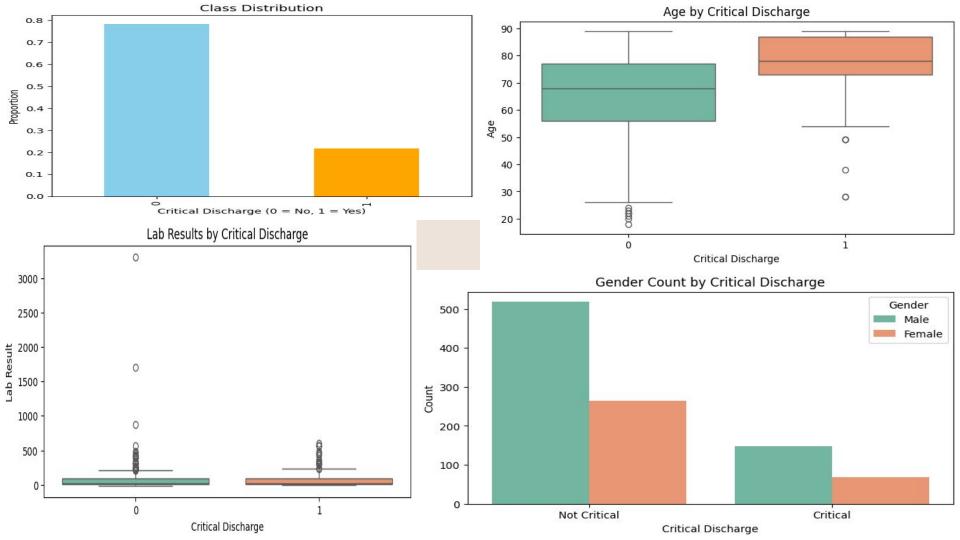
Feature Engineering & EDA

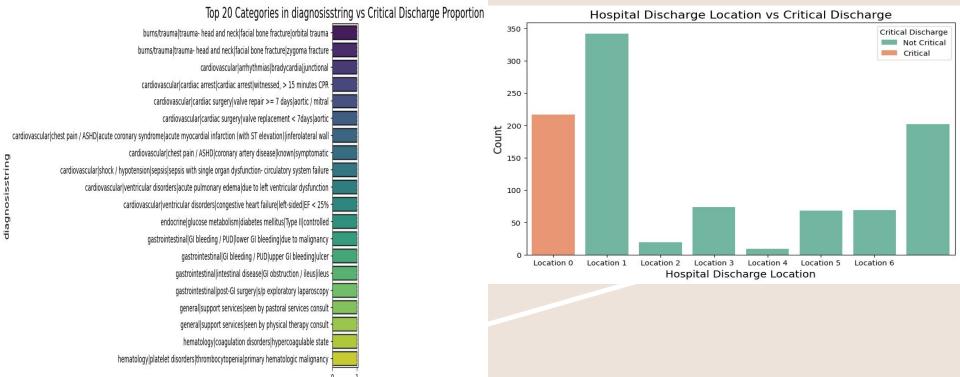
New Features Created:

• Age Groups, Diagnosis Length, Gender Encoding, Critical Discharge Indicator.

Target variable

- Research has also shown that discharge outcomes are strong indicators of readmission risk
- Binary feature
- Early identification of high-risk patients to enhance post-discharge care and reduce readmission rates.
- Imbalanced distribution (78.3% non-critical, 21.7% critical)





burns/trauma-related diagnoses) have the highest association with readmission

Proportion of Critical Discharge = 1

Model Training & Results

Best Model: Gradient Boosting Machine (GBM).

Performance Metrics:

- **AUC**: 1.0
- Log Loss: 3.459
- Optimal Threshold: 0.993 (for best F1-score).

model_id	rmse	mse	mae	rmsle	mean_residual_deviance	Classific	cation	Report:			
GBM_grid_1_AutoML_1_20250120_172333_model_17	1.39721e-07	1.9522e-14	7.27513e-08	7.05875e-08	1.9522e-14			precision	recall	f1-score	support
GBM_grid_1_AutoML_1_20250120_172333_model_19	1.39721e-07	1.9522e-14	7.27513e-08	7.05875e-08	1.9522e-14			,			
GBM_grid_1_AutoML_1_20250120_172333_model_2	1.39721e-07	1.9522e-14	7.27513e-08	7.05875e-08	1.9522e-14		0	0.99	1.00	1.00	157
GBM_grid_1_AutoML_1_20250120_172333_model_13	1.39721e-07	1.9522e-14	7.27513e-08	7.05875e-08	1.9522e-14						
GBM_grid_1_AutoML_1_20250120_172333_model_1	3.30671e-07	1.09343e-13	8.27802e-08	3.11002e-07	1.09343e-13		1	1.00	0.98	0.99	43
GBM_grid_1_AutoML_1_20250120_172333_model_16	0.000634837	4.03018e-07	0.000277034	0.000408196	4.03018e-07						
StackedEnsemble_AllModels_1_AutoML_1_20250120_172333	0.00273976	7.50627e-06	0.00156764	0.00172258	7.50627e-06	accur	acy			0.99	200
GLM_1_AutoML_1_20250120_172333	0.0034585	1.19612e-05	0.00285516	0.00282503	1.19612e-05	macro	avg	1.00	0.99	0.99	200
StackedEnsemble_BestOfFamily_1_AutoML_1_20250120_172333	0.00374225	1.40044e-05	0.00224006	0.00249188	1.40044e-05	weighted		1.00	0.99	0.99	200
GBM_2_AutoML_1_20250120_172333	0.0110038	0.000121084	0.00605146	0.00810294	0.000121084	weighten	avg	1.00	0.55	0.33	200
[30 rows x 6 columns]		Variable Imp	ortance	ς.							
AUC: 1.0						variable imp	varial		rtance scale	d_importance	percentage
						hospitaldisc	hargestat	tus 577.27	577.2741089		0.4031135
Logloss: 3.459647956244093e-05						hospitaldischa	rgelocati	on 329.38	329.3840332		0.2300106
Confusion Matrix:						diag	gnosisstri	ng 116.74	116.7477036		0.0815255
						trea	tmentstri	ng 67.14	67.1490936		0.0468906
Confusion Matrix (Act/Pred) for max f1 @ threshold = 0.9931157914848616						admitd	ctext_leng	gth 58.74	58.7481613		0.0410241
0 1 Error Rate						age_gro	0.5	53.1897964		0.0371427	
					hospitala			47.4351959		0.0331242	
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0 157 0 0 (0.0/)	157.0)						ethnic	-	48671	0.0675327 0.0580860	0.0272233
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Total 157 43 0 (0.0/)	200.0)					icd9co		23.4633026		0.0163845	
							is_fema	10.77	91300	0.0186725	0.0075271

Why GBM is the best?

Boosting Reduces Errors Iteratively:

GBM builds trees sequentially, correcting mistakes from previous iterations, making it more robust for structured medical data.

The dataset has 78.3% non-critical vs. 21.7% critical discharges.

GBM can handle imbalance effectively by assigning higher weights to minority-class samples.

Performs Well with Mixed Data Types:

H2O AutoML **tunes hyperparameters** automatically, finding the best GBM settings for your data.

Future Work

- XGBoost can be included: XGBoost requires more memory than H2O's native GBM. If AutoML detects potential memory issues, it disables XGBoost automatically.
- Check for overfitting
- Data related to expired people can be removed when working with the whole dataset as it will form a significant group. However, for this study it was not excluded since the data suggests otherwise.

(From the data we can see that some patients are alive and are not critically discharged even though they are categorized under deadly diseases while some others die from minor infections. If removed from this subset valuable information related to other variables and features will be lost. Overall, the numbers correspond to the class imbalance in the target column.)