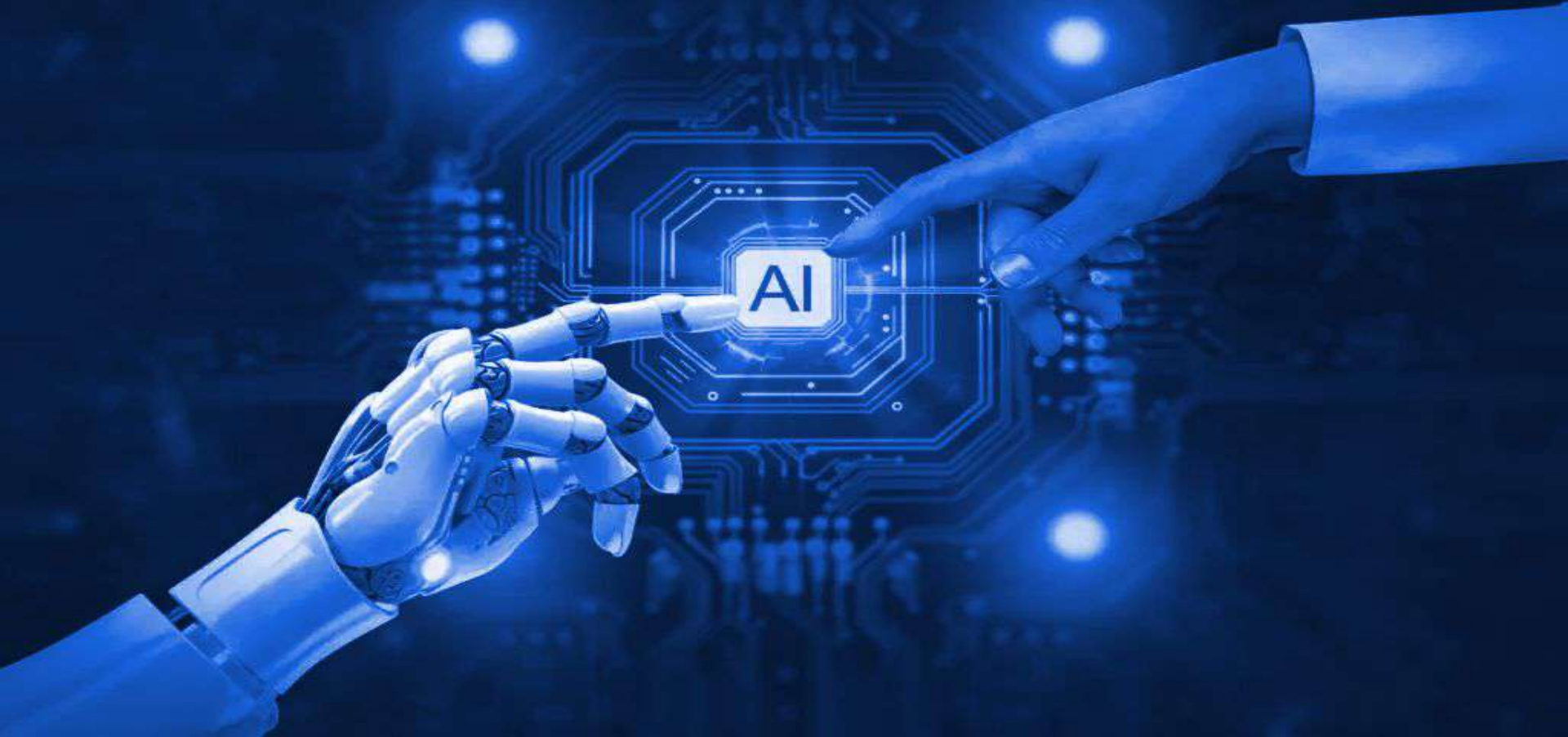
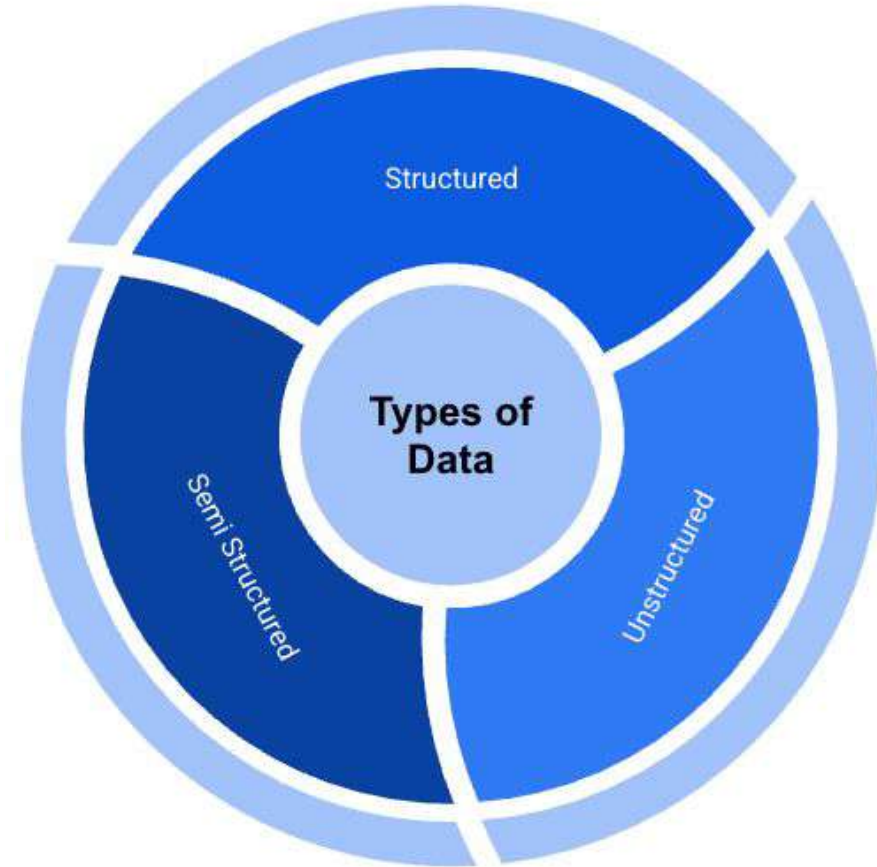


AI for Multimodal Healthcare Data

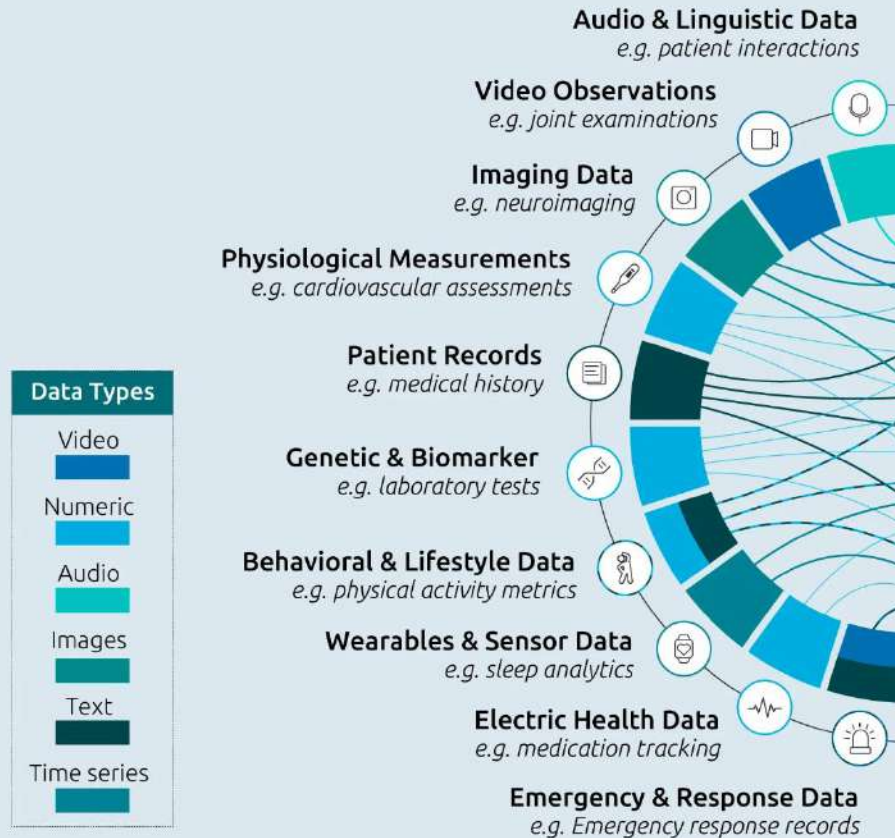


- Why AI in healthcare ?

- Types of Healthcare Data



Multimodal Data



Case Study – Shock Modes

Shock: What is it?

- A life-threatening condition where tissues don't get enough oxygen and nutrients. Leads to organ failure if untreated.

Causes of Shock

- Severe blood or fluid loss; heart fails to pump effectively; widespread infection causes blood vessel dilation + leakage.

Why Treating Early Matters

- Mortality Rates : 30 to 40%

Shock Index (SI)

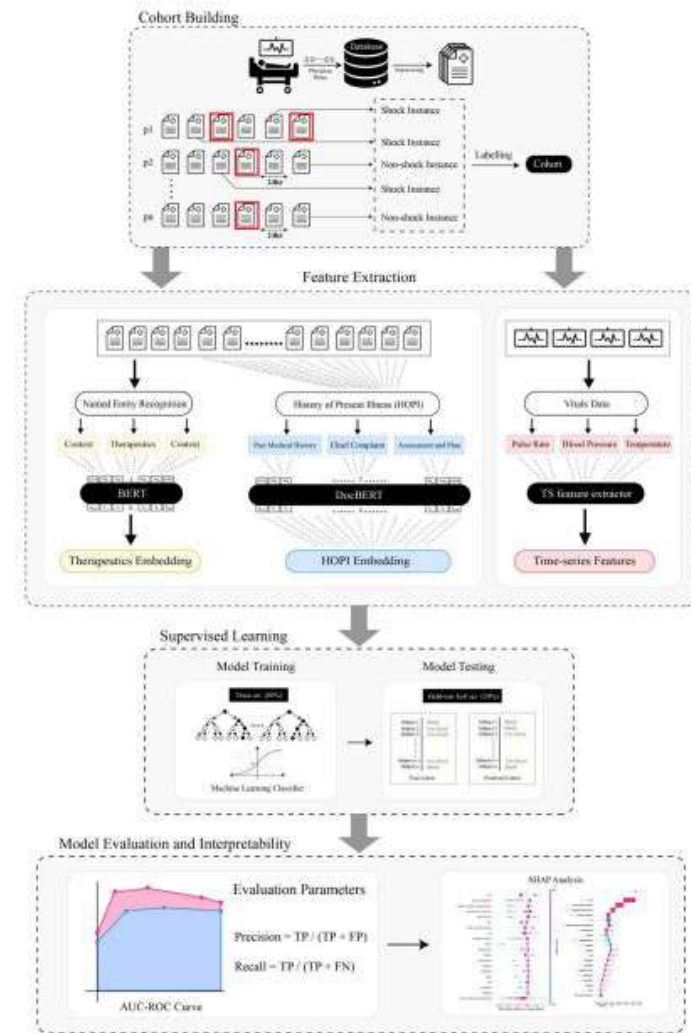
- Heart Rate (HR)/Systolic Blood Pressure (SBP)
- Abnormal: $SI \geq 0.7$ → higher risk of circulatory collapse

ShockModes Goal

- Build an Early Warning System.
- Predict abnormal SI 24 hours in advance.
- Use both vitals + physician notes.




Workflow



- Collect vitals + notes (MIMIC-III dataset).
- Clean & preprocess data.
- Extract features:
 - Vitals → time-series patterns.
 - Notes → keywords, embeddings.
- Fuse features.
- Train ML models (Random Forest, Boosting).
- Output: Risk prediction for abnormal SI.






Dataset

- Used MIMIC-III database with ICU data from 17,294 ICU-stays.
- Curated 24-hour patient encounter cohorts combining Noteevents and vital signs.
- Extracted key vitals: HR, SBP, RR, and SpO₂ for each 24-hour window.

ShareAboutExplore  

Database  Credentialed Access 

MIMIC-III Clinical Database

Alistair Johnson , Tom Pollard , Roger Mark 

Published: Sept. 4, 2016. Version: 1.4

MIMIC-III available on the cloud (Sept. 30, 2019, 2:28 p.m.)

The MIMIC-III database is now available on two major cloud platforms: Google Cloud Platform (GCP) and Amazon Web Services (AWS). To access the data on the cloud, simply add the relevant cloud identifier to your PhysioNet profile. Further instructions are available on the [MIMIC-III website](#).

Tutorials are available for [using MIMIC-III with BigQuery \(GCP\)](#) and [using MIMIC-III with Athena \(AWS\)](#).

Features from Notes

NHS Digital

SNOMED CT Browser

© SNOMED International 2017 v1.36.4 - Hosted and maintained by NHS Digital

Taxonomy

Search

Favorites

Refset

Taxonomy

Inferred view

SNOMED CT Concept

Body structure (body structure)

Clinical finding (finding)

Environment or geographical location (environment / location)

Event (event)

Observable entity (observable entity)

Organism (organism)

Pharmaceutical / biologic product (product)

Physical force (physical force)

Physical object (physical object)

Procedure (procedure)

Qualifier value (qualifier value)

Record artifact (record artifact)

Situation with explicit context (situation)

SNOMED CT Model Component (model component)

Release: United Kingdom Edition v20250730

Perspective: Full

About

SNOMED International

Leading healthcare terminology worldwide

Concept Details

Concept Details

SummaryDetailsDiagramExpressionRefsetsMembersReferencesClassification Map

Stated

Inferred

Parents

SNOMED CT Concept (SNOMED RT+CTV3)

Clinical finding (finding)

SCTID: 404684003

404684003 | Clinical finding (finding) |

Clinical finding (finding)

Clinical finding

No attributes

Features from Vitals

- Extracted 3117 time-series features using tsfresh.

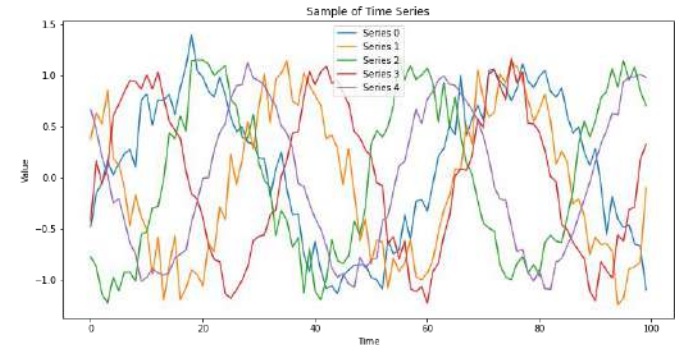
- Examples:

Autocorrelation (patterns)

Fourier transforms (signal changes)

Entropy (randomness in signals)

- tsfresh converts raw time-series into thousands of features (mean, variance, entropy, Fourier coefficients, etc.).
- Extracted features can be fed into ML models for tasks like classification, regression, clustering, or forecasting.



Model Development and Validation

- Input features: vitals (tsfresh), embeddings of therapeutics, and HOPI.
- Feature reduction: Used Extra Trees classifier (scikit-learn) to select important features.
- Imbalanced data: Handled using SMOTE oversampling (scikit-learn).
- Models used: Logistic Regression, Random Forest, GradientBoost, AdaBoost, XGBoost.
- Evaluation: Performed with bootstrap sampling (100 iterations); key metrics were AUC-ROC and F1-score.

SHAP Analysis for Interpretability

- SHAP (Shapley Additive Explanations): Uses Shapley values to calculate feature importance and explain model outputs.
- Interpretability method: Prior probability from training data is compared with model output probability to explain predictions.
- Local explanations: Patient-specific predictions visualized using waterfall plots, showing which features pushed toward normal/abnormal SI.
- Global explanations: Overall feature importance visualized with bar plots and beeswarm plots.

