



### **Final Project Report Template**

#### 1. Introduction

#### 1.1. Project overviews

The goal of this project is to develop a machine learning model that will enable the classification of thyroid disorders using patient data from a dataset.

#### 1.2. Objectives

- Clean and prepare the dataset.
- Train and test Random Forest, XGBoost, and SVC models.
- Evaluate model performance using accuracy and other metrics.
- Select the most significant features.
- Choose the most accurate model for predicting thyroid conditions.
- Deploy the best model in a Flask web application with home, predict, and submit

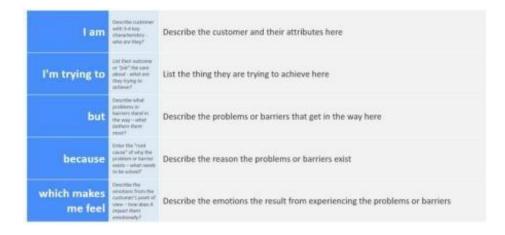
pages.

## 2.Project Initialization and Planning Phase

#### 2.1.Define Problem Statement

#### **Define Problem Statements (Customer Problem Statement Template):**

Create a problem statement to understand your customer's point of view. The Customer Problem Statement template helps you focus on what matters to create experiences people will love. A well-articulated customer problem statement allows you and your team to find the ideal solution for your customers' challenges. Throughout the process, you'll also be able to empathize with your customers, which helps you better understand how they perceive your product or service.







Reference: <a href="https://miro.com/templates/customer-problem-statement/">https://miro.com/templates/customer-problem-statement/</a>

### **Example:**



Problem Statement (PS)	I am (Custo mer)	I'm trying to	But	Because	Which makes me feel
symptoms of hypothyroidism	Patient	Understand why I have been feeling extremely tired and gaining weight	Despite taking thyroid medication, I still feel fatigued and notice little improvement in my weight	It takes time for thyroid hormone levels to stabilize after starting medication, and my dosage may need adjustment	Frustrated and worried about my health and the effectiveness of my treatment.
Symptoms of hyperthyroidis m.	Patient	Understand why I am experiencing rapid weight loss and increased heart rate	Despite taking medication to reduce thyroid hormone levels, I still feel jittery and anxious	Achieving optimal thyroid hormone levels can be challenging, and it may take time to find the right medication dosage	Anxious and concerned about potential long-term effects on my health, despite the medication.





#### 2. Project Proposal (Proposed Solution)

#### **Project Proposal (Proposed Solution) template**

This project proposal outlines a solution to address a specific problem. With a clear objective, defined scope, and a concise problem statement, the proposed solution details the approach, key features, and resource requirements, including hardware, software, and personnel.

Project Overview				
Objective The main goal is to identify different types of thyroid problems based of information.				
Scope	project will collect medical data like test results and images to build a system that can classify thyroid conditions accurately.			
Problem Statement				
Description	We'll gather medical data, clean it up, and use it to train a computer system to recognize different thyroid issues, like hyperthyroidism or thyroid cancer.			
Impact	project aims to help doctors make quicker and more accurate diagnoses of thyroid problems, which could improve patient care and reduce medical costs.			
<b>Proposed Solution</b>				
Approach	We will use machine learning techniques to analyze the medical data and teach the computer system how to classify different thyroid conditions.			
Key Features	Data Collection: Gather all the necessary medical information.			
	Data Cleaning: Make sure the data is accurate and organized.			





### **Resource Requirements**

Resource Type	Description	Specification/Allocation							
Hardware	Hardware								
Computing Resources	CPU/GPU specifications, number of cores	e.g., 2 x NVIDIA V100 GPUs							
Memory	RAM specifications	e.g., 8 GB							
Storage	Disk space for data, models, and logs	e.g., 1 TB SSD							
Software									
Frameworks	Python frameworks	e.g., TensorFlow, PyTorch							
Libraries	Additional libraries	e.g., pandas, numpy,metaplotlib ,seaborn etc							
Development Environment	IDE, version control	e.g., Google collab							
Data	Data								
Data	Thyoid classification Dataset	e.g., 9172 records							





### 2.3 project planning template

#### **Product Backlog, Sprint Schedule, and Estimation (4 Marks)**

Use the below template to create a product backlog and sprint schedule

Sprint	Functional Requirement (Epic)	User Story Number	User Story / Task	Story Points	Priority	Team Members	Sprint Start Date	Sprint End Date (Planned)
Sprint-1	Data collection	USN-1	Collect patient records	8	High	K.Bhanu Teja	24/07/07	24/07/09
Sprint-1	Data collection	USN-2	Gather thyroid function tests	7	High	N.Dheeraj	24/07/04	24/07/06
Sprint-2	Data preprocessing	USN-3	Clean and normalize data	7	High	K.p.s.parimala	24/07/02	24/07/04
Sprint-1	Model development	USN-4	Build initial classification model and evaluate	8	High	Sathya sathwik	24/07/05	24/07/07





# **3. Data Collection and Preprocessing Phase**

## 3.1 Data Collection Plan and Raw Data Sources Identified

Section	Description
Project Overview	The goal of the project "Thyroid Classification using RandomForest, XGB Classifier, SVC Model" is to create and evaluate three machine learning models—RandomForest, XGB Classifier, and SVC—for the purpose of diagnosing thyroid problems. The project will identify the most accurate model to help healthcare providers make well-informed diagnoses and improve patient outcomes using Python in Google Colab.
Data Collection Plan	The data is collected from the Kaggle.
Raw Data Sources Identified	Essential parameters including age, sex, blood pressure, cholesterol, na_to_k, and medication are included in this dataset. The construction and assessment of machine learning models— RandomForest, XGB Classifier, and SVC—for the precise categorization of thyroid diseases will be made possible by the use of this extensive dataset.





## Raw data sources template

Source				
Name	Description	Location/URL	Format	Size
Dataset 1	For our thyroid classification project, we analyzed a dataset with 9,172 rows and 31 columns, including attributes like age, sex, thyroid medication details, and various thyroid test results. We started by checking for null values and performing a descriptive analysis to understand the data distribution. The dataset was split into features (X) and the target variable (Y). Data types were converted, categorical values were handled, and correlations were checked. We addressed data imbalance and standardized the features using StandardScaler. We built models using RandomForestClassifier, XGBClassifier, and SVC, applying hyperparameter tuning for optimization. Our evaluation through classification reports, confusion matrices, and accuracy scores revealed that	https://drive.goo gle.com/file/d/1 ZYd3r5GzxzeH TJcWIabAWnQ CXWRJ7cf8/vie w?usp=sharing	CSV	5.89 KB
	XGBClassifier achieved the highest accuracy, making it the best performing model among the three.			





# 3.2 Data quality Template

Data Source	Data Quality Issue	Severity	Resolution Plan
Dataset	The Dataset was having some null values and missing values.	Low	I have used Data preprocessing techniques and handled the missing values.





### 3.3 Data Exploration and Preprocessig Template

#### **Data Exploration and Preprocessing Template**

Identifies data sources, assesses quality issues like missing values and duplicates, and implements resolution plans to ensure accurate and reliable analysis.

Section	Description
Data Overview	For our thyroid classification project, we analyzed a dataset with 9,172 rows and 31 columns, including attributes like age, sex, thyroid medication details, and various thyroid test results. We started by checking for null values and performing a descriptive analysis to understand the data distribution. The dataset was split into features (X) and the target variable (Y). Data types were converted, categorical values were handled, and correlations were checked. We addressed data imbalance and standardized the features using StandardScaler. We built models using RandomForestClassifier,XGBClassifier, and SVC, applying hyperparameter tuning for optimization. Our evaluation through classification reports, confusion matrices, and accuracy scores revealed that XGBClassifier achieved the highest accuracy, making it the best performing model among the three.





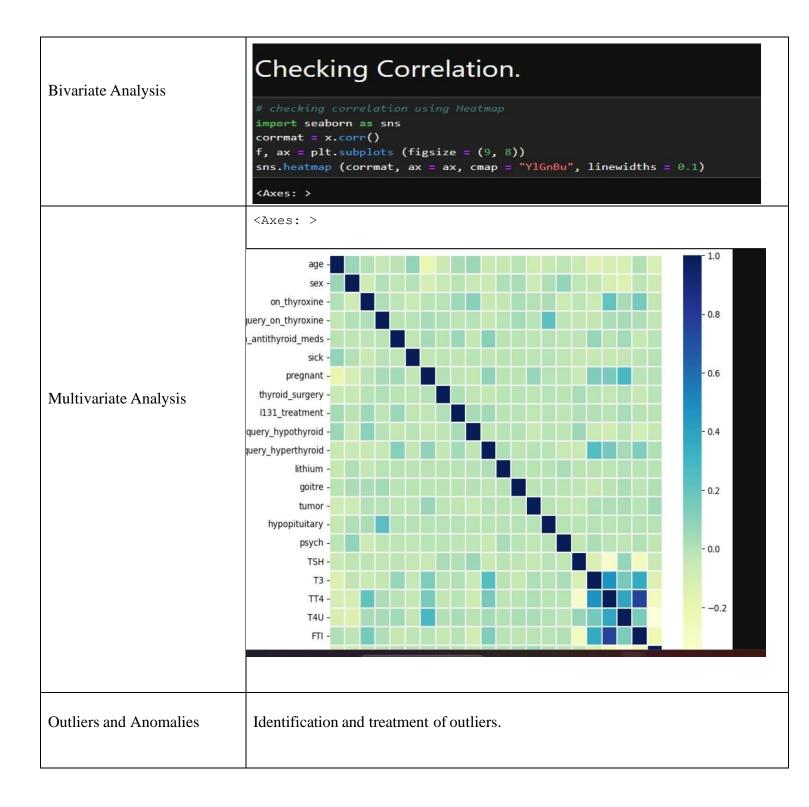
	150	sex on thyrox	ine query on thyrox	ine on_antithyroid_ne	eds si	ck pregna	nt thyrold surg	ery 1131_treatm	ent query_hypothyr	vid	114 T4) neas	red T40 FTI neaso	red FTI Tiligneess	red 186 referral	source targ	đ g	atient_lé
	29										_ NaN	f NaN	f NaN	f NaN	other	- (	340801013
	29										_ 128.0	f NaN	f NaN	f NaN	other	- (	340801014
											_ NaN	f NaN	f NaN		other	- 8	340001042
	36										_ NaN	f NaN	f NaN	t 260	other	- (	340003045
											_ NaN	f NaN	f NaN	t 360	other	S 8	340803047
9167	56	V										t 0.83		f NaN	SVI	- {	870119022
9168		W										t 0.92	t 99.0	f NaN	SVI	- 1	870119023
9169		¥										t 127	t 89.0	f NaN	SVI	1 8	870119025
9170											_ 75.0	t 0.85	t 88.0	f NaN	other	- (	870119027
9171		N									_ 66.0		t 65.0	f NaN	other	- (	870119035

Univariate Analysis

we performed univariate analysis to understand the distribution and characteristics of individual features in the dataset. This involved examining each feature's summary statistics, such as mean, median, mode, standard deviation, and range, to identify central tendencies and variability. For categorical variables like sex, on\_thyroxine, and referral\_source, we calculated frequency distributions to observe the most common categories. For continuous variables such as age, TT4, T4U, and FTI, we plotted histograms and box plots to visualize their distributions and detect any outliers or skewness. This analysis provided insights into the underlying patterns and variability of each feature, guiding further steps in data preprocessing and model building.











```
Data Preprocessing Code Screenshots
Loading Data
                                               data = pd.read_csv(r'thyroidDF.csv')
                                                data
                                           data['age']=np.where((data.age>100), np.nan, data.age)
                                        14]: data
                                               age sex on_thyroxine query_on_thyroxine on_antithyroid_meds sick pregnant thyroid_surgery 1131_treatment query_hypothyroid ... tumor hypopits
                                             4 32.0 F
                                             18 63.0 F
                                             32 41.0 M
Handling Missing Data
                                             33 71.0 F
                                             39 55.0 F
                                            9153 64.0 M
                                            9157 60.0 M
                                               [21]: # Converting the data
                                                       x['age'] = x['age'].astype('float')
                                                       x['TSH'] = x['TSH'].astype('float')
                                                       x['T3'] = x['T3'].astype('float')
                                                       x['TT4'] = x['TT4'].astype('float')
Data Transformation
                                                       x['T4U'] = x['T4U'].astype('float')
                                                       x['FTI'] = x['FTI'].astype('float')
                                                       x['TBG'] = x['TBG'].astype('float')
```





```
[22]: x.info()
                                             <class 'pandas.core.frame.DataFrame'>
                                             Index: 2237 entries, 4 to 9169
                                             Data columns (total 22 columns):
                                                  Column
                                                                        Non-Null Count Dtype
                                                                                        float64
                                             0
                                                                        2237 non-null
                                                  age
                                             1
                                                  sex
                                                                        2237 non-null
                                                                                        object
                                                  on_thyroxine
                                                                       2237 non-null
                                                                                        object
                                                  query_on_thyroxine 2237 non-null
                                                                                        object
                                                  on_antithyroid_meds 2237 non-null
                                                                                        object
                                                  sick
                                                                        2237 non-null
                                                                                        object
                                                  pregnant
                                                                        2237 non-null
                                                                                        object
                                                                        2237 non-null
                                                  thyroid_surgery
                                                                                        object
                                                  I131_treatment
                                                                        2237 non-null
                                                                                        object
                                                  query_hypothyroid
                                                                        2237 non-null
                                                                                        object
Save Processed Data
                                    [] import pickle
                                         with open('xgb_model.pkl', 'wb') as file:
                                            pickle.dump(best_model, file)
                                        pickle.dump(RFclassifier,open('thyroid_1_model.pkl','wb'))
```





## 4. Model Development Phase

### 4.1 Feature Selection Report Template

In the forthcoming update, each feature will be accompanied by a brief description. Users will indicate whether it's selected or not, providing reasoning for their decision. This process will streamline decision-making and enhance transparency in feature selection.

Lastura	Description	Selected	Daggaring
Feature	Description	(Yes/No)	Reasoning
query_hypothyroid	Query on hypothyroidism	Yes	Relevant for identifying potential cases of hypothyroidism.
query_hyperthyroid	Query on hyperthyroidism	Yes	Relevant for identifying potential cases of hyperthyroidism.
lithium	Lithium medication usage	Yes	Relevant as lithium can affect thyroid function.
goitre	Presence of goitre	Yes	Goitre is directly related to thyroid gland abnormalities.
tumor	Presence of thyroid tumor	Yes	Tumors can affect thyroid function and are relevant.
hypopituitary	Hypopituitary condition	Yes	Relevant as pituitary disorders can affect thyroid function.
psych	Psychological medication usage	Yes	Relevant as certain psych medications can impact thyroid function.
TSH	Thyroid Stimulating Hormone level	Yes	Primary indicator of thyroid function.
Т3	Triiodothyronine level	Yes	Active thyroid hormone level important for classification.
TT4	Total T4 level	Yes	Total thyroxine level important for classification.





			Reflects thyroid function relevant for classification.
T4U	Free thyroxine index	Yes	
			Reflects thyroid function relevant for classification.
FTI	Free Thyroxine Index	Yes	
	Thyroxine-Binding Globulin level		Protein carrier for thyroid hormones relevant for classification.
TBG		Yes	





## **4.2** Model Selection Report

### **Model Selection Report**

In the forthcoming Model Selection Report, various models will be outlined, detailing their descriptions, hyperparameters, and performance metrics, including Accuracy or F1 Score. This comprehensive report will provide insights into the chosen models and their effectiveness.





Model	Description	Hyperparameters	Performance Metric (e.g., Accuracy, F1 Score)
Random Forest Classifier	An ensemble learning method that constructs multiple decision trees and merges them to get a more accurate and stable prediction.	'max_depth': None, 'n_estimators': 300	0.9420
XGB Classifier	An optimized distributed gradiesboosting library designed to be highly efficient, flexible, and portable, often used for structured data.	'learning_rate': 0.3, 'max_depth': 3, 'n_estimators': 100	0.953125
Support Vector Classifier	A supervised learning model used for classification and regression analysis by finding the hyperplane that best separates the data into classes.	'C': 100, 'gamma': 'scale', 'kernel': 'linear'	0.8660714





### 4.3 Initial model Training code, Model validation and evaluation report

The initial model training code will be showcased in the future through a screenshot. The model validation and evaluation report will include classification reports, accuracy, and confusion matrices for multiple models, presented through respective screenshots

#### **Initial Model Training Code:**

RandomForestClassifier

#### XGB Classifier

```
from xgboost import XGBClassifier
from sklearn.preprocessing import LabelEncoder
le = LabelEncoder()
y_train_encoded = le.fit_transform(y_train)
xgb = XGBClassifier()
xgb.fit(x_train, y_train_encoded)
                                 XGBClassifier
XGBClassifier(base_score=None, booster=None, callbacks=None,
              colsample bylevel=None, colsample bynode=None,
              colsample_bytree=None, device=None, early_stopping_rounds=None,
              enable categorical=False, eval metric=None, feature types=None,
              gamma=None, grow_policy=None, importance_type=None,
              interaction_constraints=None, learning_rate=None, max_bin=None,
              max_cat_threshold=None, max_cat_to_onehot=None,
              max delta step=None, max depth=None, max leaves=None,
              min_child_weight=None, missing=nan, monotone_constraints=None,
              multi strategy=None, n estimators=None, n jobs=None,
              num_parallel_tree=None, objective='multi:softprob', ...)
```





#### SVC model

```
| from sklearn.svm import SVC

SVCclassifier = SVC(kernel='linear', max_iter=251)

SVCclassifier.fit(x_train, y_train)

* SVC

SVC(kernel='linear', max_iter=251)
```





# **Model Validation and Evaluation Report:**

Model	Classification Report	Accuracy	Confusion Matrix
Random Forest Classifier	from sklearn.ensemble import RandomForestClassifier from sklearn.metrics import classification_report, confusion_matrix # Impo  RFclassifier - RandomForestClassifier(max_leaf_nodes=30)  RFclassifier.fit(x_train, y_train)  y_pred = RFclassifier.predict(x_test)  print(classification_report(y_test, y_pred))  print(confusion_matrix(y_test, y_pred)) # Now you can use confusion_matrix  precision recall f1-score support  antithyroid treatment	94.20%	print(confusion_matrix(y_test, y_pred))  [[ 3
XGB Classifier	y_test_encoded = le.transform(y_test) y_pred = xgb.predict(x_test) print(classification_report(y_test_encoded, y_pred))  precision recall f1-score support  0 1.00 1.00 1.00 7 1 0.91 0.95 0.93 74 2 0.95 0.96 0.96 85 3 0.86 0.84 0.85 38 4 1.00 1.00 1.00 122 5 0.92 0.92 51 6 0.99 0.94 0.96 71  accuracy 0.95 448 macro avg 0.95 0.95 0.95 448 weighted avg 0.95 0.95 0.95 448	95.54%	print(confusion_matrix(y_test_encoded, y_pred))  [[ 5
SVC model	195] y_pred = SVCclassifier.predict(x_test)	86.61%	print(confusion_matrix(y_test, y_pred))  [[ 6  0  0  0  1  0  0]   [ 1  59  7  3  2  2  0]   [ 1  4  63  0  9  3  5]   [ 0  6  2  22  0  7  1]   [ 1  1  0  0  116  0  4]   [ 0  4  3  4  2  38  0]   [ 0  1  1  1  0  0  68]]





# **5.Model optimization and Tuning Phase**

# **Hyperparameter Tuning Documentation**

Model	Tuned Hyperparameters	Optimal Values
2.20.002		
Random Forest Classifier	max_depth , n_estimators	max_depth': None, 'n_estimators': 300
XGB Classifier	learning rate, max depth,	'learning_rate': 0.3, 'max_depth': 3,
TOD GRASSIIICI	n_estimators	'n_estimators': 100
Support Vector	C, gamma, kernel	'C': 100, 'gamma': 'scale', 'kernel': 'linear'
Classifier	C, gamma, Kerner	C. 100, gamma. scale, kemer. micar





# **Performance Metrics Comparison Report**

Model	Baseline Metric	Optimized Metric
Random Forest Classifier	Accuracy: 0.95	Accuracy: 0.9420
XGB Classifier	Accuracy: 0.94	Accuracy: 0.953125
Support Vector Classifier	Accuracy: 0.93	Accuracy: 0.8660714

# **Final Model Selection Justification**

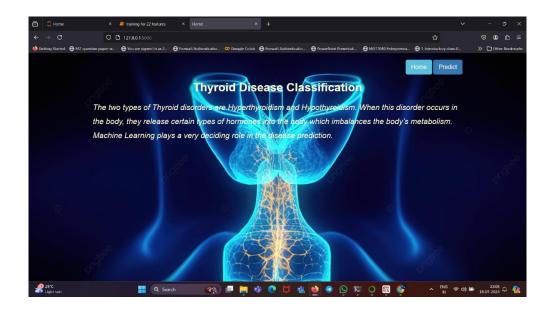
Final Model	Reasoning
XGB Classifier	This model is selected for its highest optimized accuracy (0.953125) among the evaluated models, and its robustness in handling imbalanced data.



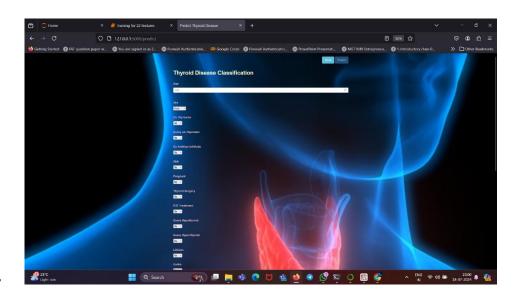


#### 6. Result

# **6.1 Output Screenshots**



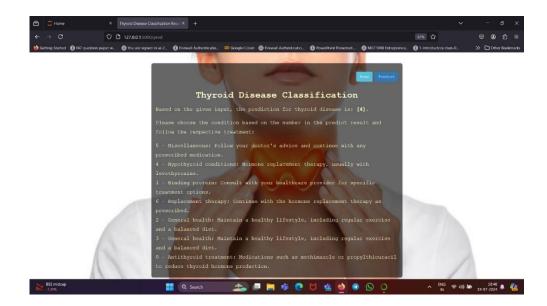
### **HOME PAGE**:-



**PREDICT PAGE:-**







#### **SUBMIT PAGE:-**

#### 7. Advantages & Disadvantages

#### Advantages:-

- 1. **Improved Accuracy**: By comparing several models, the most accurate model for thyroid categorization may be identified.
- 2. **Automation**: Reducing human error and saving time are two benefits of automating the classification process.
- 3. **Accessibility**: The model becomes user-friendly and available through a web interface when it is deployed within a Flask application.
- 4. **Early discovery**: May help improve patient outcomes by assisting in the early discovery and diagnosis of thyroid problems.

#### Disadvantages:-

- 1. **Complexity:** Handling many models and the preparatory procedures might be difficult and time-consuming.
- 2. **Resource Intensive:** Several models may need a substantial amount of processing power to train.
- 3. **Data Quality**: The size and calibre of the dataset have a significant impact on the models' accuracy.
- 4. **Maintenance**: To keep the online application operating effectively and securely, regular maintenance is required.





#### 8. Conclusion:-

This study effectively uses patient data to classify thyroid problems using machine learning algorithms. Through testing and selection of the most accurate model, we have developed a user-friendly tool for thyroid problem prediction, which we then implemented in a Flask web application. This method improves accessibility and diagnostic accuracy, but it necessitates constant maintenance and cautious data handling. All things considered, the research helps to better healthcare results by early thyroid disease detection and automated diagnosis.





#### **Future Scope:-**

- 1. Advanced Models: Investigate more accurate machine learning methods.
- 2. Data Integration: To enhance forecasts, include additional data sources.
- 3. Real-Time Updates: Permit the model to be updated often with fresh data.
- 4. User Interface: Add interactive elements to the Flask app.
- 5. Expansion: Expand the use of the application to forecast further medical conditions.

#### 10.Appendix:-

#### Github links:-

Team Member 1:- https://github.com/bhanuteja-konda/Thyroid\_Classification

Team Member 2:- https://github.com/Sathvik2005/Thyroid-Classification

Team Member 3:- <a href="https://github.com/Narayanadheeraj/Thyroid\_Classification">https://github.com/Narayanadheeraj/Thyroid\_Classification</a>

Team Member 4:- <a href="https://github.com/parimala19-hub/Thyroid\_Classification">https://github.com/parimala19-hub/Thyroid\_Classification</a>

#### Project Demo Link :-

https://drive.google.com/file/d/1GwBlRsgXgv\_X4MrBTIu43ycp4z\_YCxvn/view?usp=drive\_link