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Thyroid Cancer Risk Analysis

Providing an in-depth analysis of dataset and uncovering key insights into which demographics are at high risk of having thyroid cancer.



Overview

Providing an in-depth analysis of dataset and uncovering key insights into which demographics are at risk of having thyroid cancer:

- **Hypothesis 1**: Age and gender influence thyroid cancer risk, with older individuals and females having a higher probability
 - Validation: Use box plots and regression analysis to explore how cancer risk varies across different age groups and gender distributions
- **Hypothesis 2**: Certain countries and ethnicities have a higher prevalence of thyroid cancer due to genetic and environmental factors
 - Validation: Conduct geospatial analysis and visualize the distribution of thyroid cancer cases across different regions.



Planning & Design

Ideation

Project Goal: Build interactive dashboards for data analysis

Medical Use Case: Improve decision-making through data insights of thyroid cancer risk based on demographics

Target Audience: Medical professionals, analysts, WHO and decision-makers

Design

User Stories: "As a Medical practitioner, we want to explore demographics of thyroid cancer risk dynamically."

Intuitive UI: Clean layouts, easy navigation

Accessibility: Readable colours, labeling

Interactivity: Clickable filters, zoomable charts and maps

Hypothesis: Which demographic has high risk of thyroid cancer

Technologies

Tools: Visual Studio Code, Jupyter Notebook, Power BI, PowerPoint

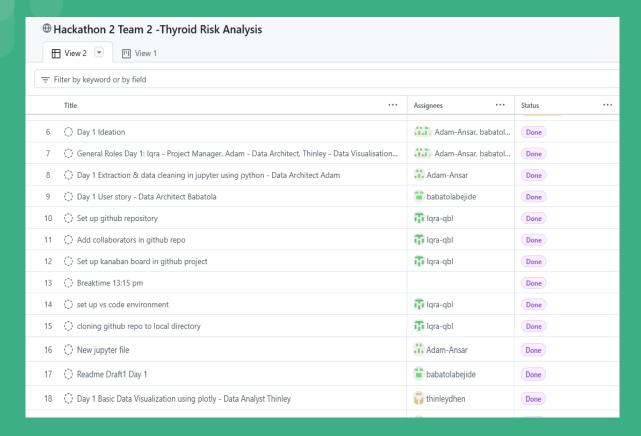
Wireframing: Balsamic Wireframes

Project Management: GitHub Projects, Google Meets

Version Control: GitHub for collaboration

Libraries & Frameworks:
Python (Pandas, NumPy, Plotly,
Seaborn, Ipywidgets,
StatsModels), Power BI

Project Board



Assigned Tasks: 30

MoSCoW Prioritisation:

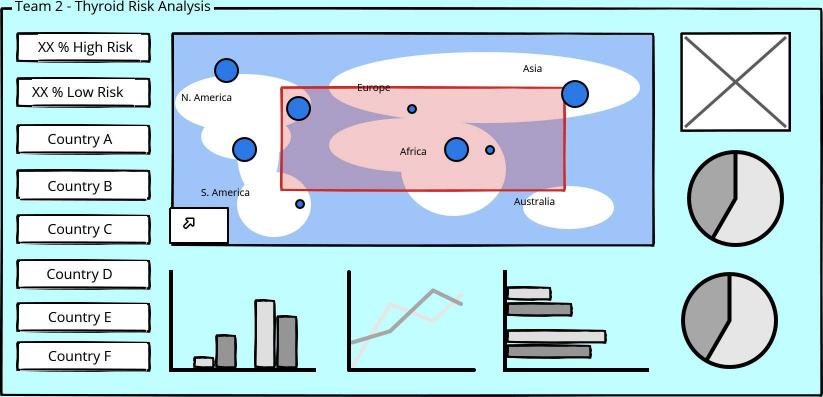
Must have: an interactive dashboard and proper documentation

Should have: simple and followable code

Could have: nicely stylized code and dashboard

Would have: utilize the dataset to full potential

Project Backlog: Setting VS Code and Github collaboration

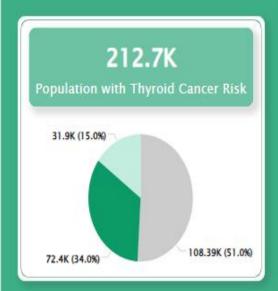


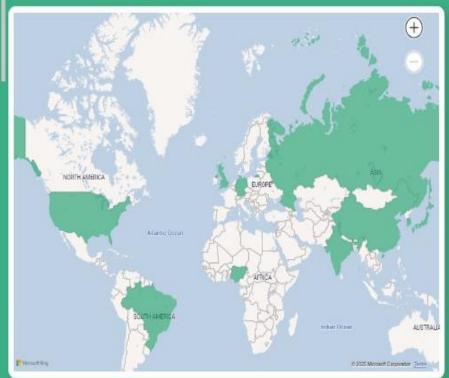
Features

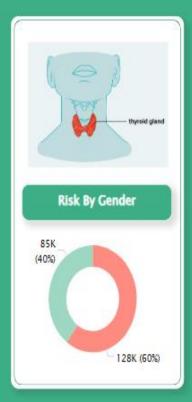
- 1. Interactive: all variables i.e. age, gender, country, ethnicity, diagnosis are interlinked so selecting one will show all the relevant data across different charts and map
- 2. At the bottom there are instructions on using the different features of the dashboard
- 3. Hovering over any chart will further explain the data specific e.g. regarding gender, amount of population at risk, country etc

Thyroid Cancer Risk Analysis 21.4K ulation with Thyroid Cancer Rit The Thyroid Cancer Risk Analysis is an interactive dashboard that allows users to select any variable to see corresponding changes in all the charts and plas. It

Thyroid Cancer Risk Analysis









Risk by Family History

Risk by Age

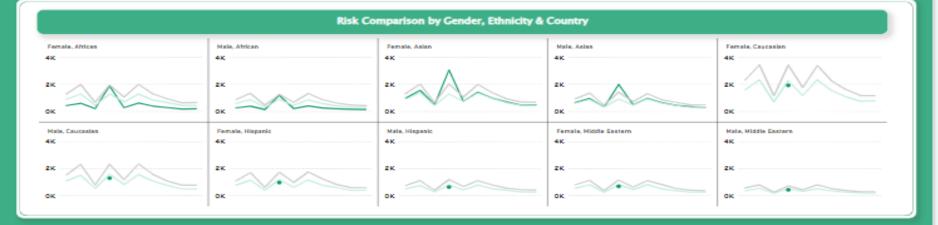
Diagnose by Gender

200K

1979

00K



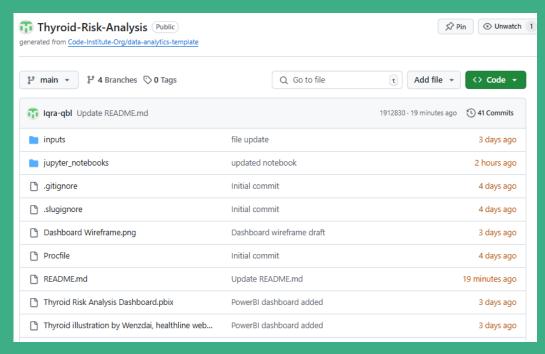


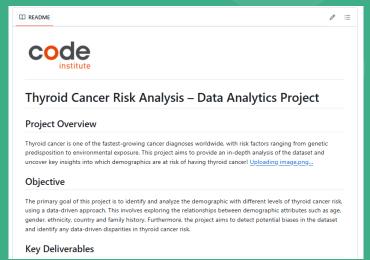
Dashboard Instructions & Information

The Thyroid Cancer Risk Analysis is an interactive dashboard that allows users to select any variable to see corresponding changes in all the charts and pies. It allows users to see what are the risk levels and diagnoses for getting thyroid cancer by gender, different age groups, ethnicity, countries etc.

To use the dashboard it is recommended to select or hover over any part or point of the charts and pies in the dashboard. For example selecting any gender in the top right pie will showcase the data associated with that gender. Similarly, selecting one of the slices of the left pie chart at top will allow the user to see the percentage of population with three different levels of thyroid cancer risk: low, medium and high; and all the data related like what country or age or ethnicity they belong to.

Documentation, Testing & Version Control





GitHub Repository: https://github.com/lqra-qbl/Thyroid-Risk-Analysis

Section 1: Data Extraction, Transformation, and Loading (ETL)

Setting up & Importing Python packages that we will be using in this project to carry out the analysis. For example Numpy to compute numerical operations and handle arrays, Pandas for data manipulation and analysis, Matplotlib, Seaborn and Plotly to create different data visualisations.

```
import pandas as pd
import numpy as np
import matplotlib.pyplot as plt
import seaborn as sns
sns.set_style('whitegrid')
import plotly.express as px
✓ 3.8s
```

Data Extraction

Loading the CSV dataset containing the data collected previously and extracting it into dataframe using pd.read_csv() function.

```
df = pd.read_csv("thyroid cancer risk data.csv")
```

Section 2: Bias Detection

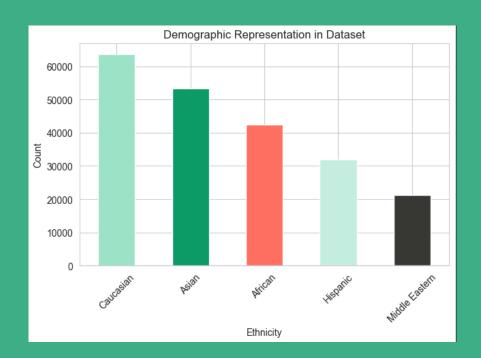
Checking the data for any bias in any of the classes

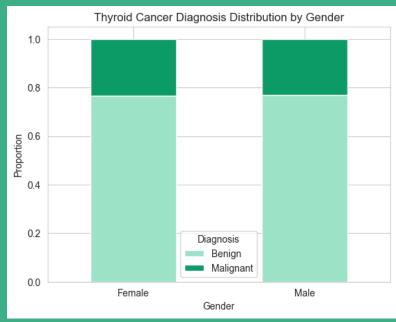
Bias in data can significantly affect the outcomes and interpretations of our analysis. It is crucial to identify and address any biases to ensure the validity and fairness of our results. In this section, we will check for potential biases in the dataset across different classes such as gender, age, ethnicity, and family history.

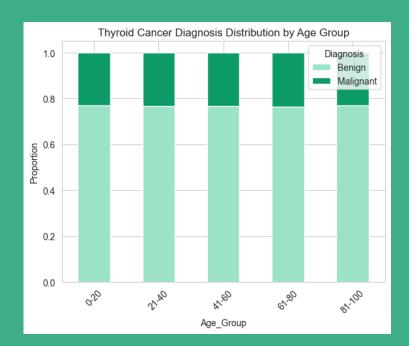
Load the cleaned dataset for Bias Analysis

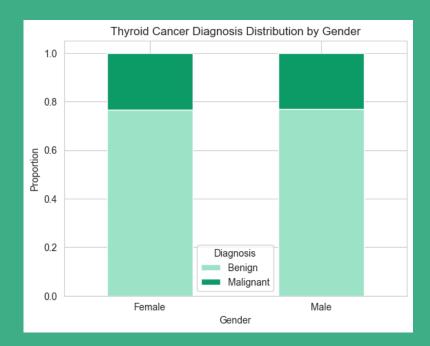
```
# Analyze demographic representation
data = pd.read_csv('thyroid_cancer_risk_data_cleaned.csv')

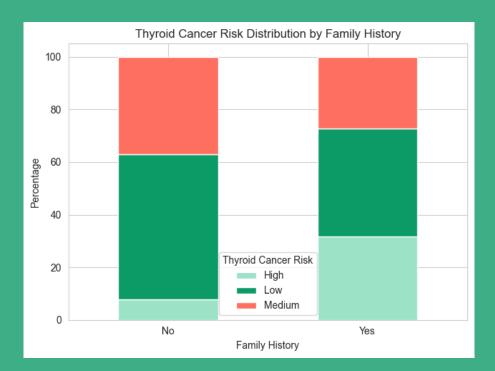
demographic_counts = data['Ethnicity'].value_counts()
print(demographic_counts)
```











Section 3: Analysis

Hypothesis 1: Age and gender influence thyroid cancer risk, with older individuals and females having a higher probability.

Validation: Use box plots and regression analysis to explore how cancer risk varies across different age groups and gender distributions. **Logistic Regression Modeling:** Fitting a logistic regression model to quantify the effect of age and gender on thyroid cancer probability.

```
Empty markdown cell, double-click or press enter to edit.
```

```
# Define independent variables (without intercept)
```

df['Thyroid Cancer Risk Numeric'] = df['Thyroid Cancer Risk'].map({'Low': 1, 'Medium': 2, 'High': 3})

from statsmodels.miscmodels.ordinal model import OrderedModel

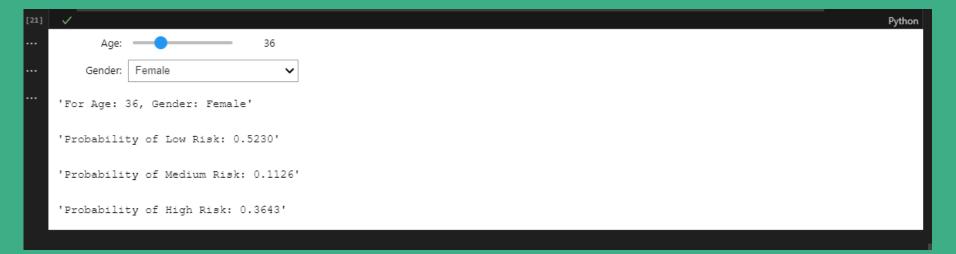
df['Gender Binary'] = df['Gender'].map({'Female': 1, 'Male': 0})

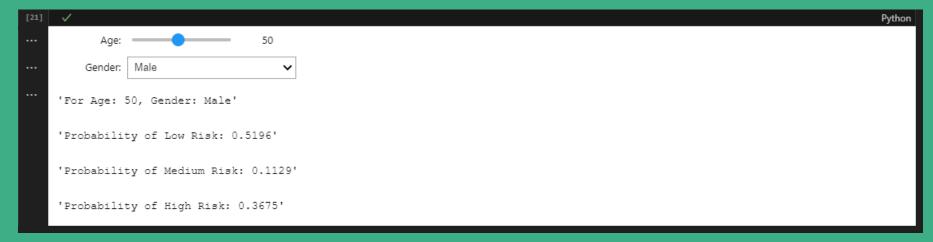
Convert categorical variables to numerical values

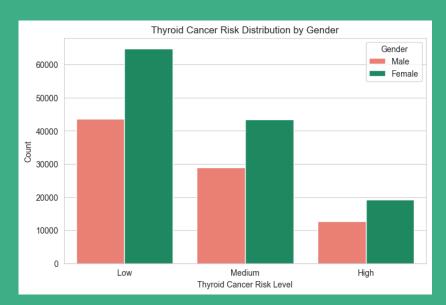
X_ordinal = df[['Age', 'Gender_Binary']]

```
Ln 14, Col 96 Spaces: 4 Spaces: 4 CRLF Cell 49 of 74 🔠 ✓ Prettier
```

```
Optimization terminated successfully.
       Current function value: 0.994911
       Iterations: 13
       Function evaluations: 16
       Gradient evaluations: 16
                           OrderedModel Results
               Thyroid Cancer Risk Numeric Log-Likelihood: -2.1161e+05
Dep. Variable:
Model:
                           OrderedModel AIC:
                                                               4.232e+05
                      Maximum Likelihood
Method:
                                       BIC:
                                                               4.233e+05
Date:
                        Mon, 10 Feb 2025
Time:
                               20:18:29
No. Observations:
                                212691
Df Residuals:
                                212687
Df Model:
                      std err z P>|z|
                                                   [0.025
                coef
                                                             0.975]
         0.0003 0.000 1.504 0.133 -8.72e-05
                                                             0.001
Age
Gender Binary 0.0178 0.008 2.111 0.035 0.001 0.034
1/2
    0.0640 0.012 5.354 0.000 0.041
                                                             0.087
2/3
              0.5284
                        0.003 155.718
                                          0.000
                                                    0.522
                                                             0.535
```









Insights & Finding

Key Data Insights

- Caucasians are the most represented group, with 63,669 entries, while the Middle Eastern individuals are the least represented, with only 21,335 entries, Asians with 53,261 entries, Africans with 42,414 entries, and Hispanics with 32,012 entries, showing an uneven distribution
- Females are more likely to have thyroid cancer especially Asian females
- Chi-square test p-value = 0.5102 (p > 0.05), which means no statistically significant difference in the diagnosis proportions across the genders

- Chi-squared test p-value for age groups: 0.5586, analysis does not show any bias in thyroid cancer diagnosis based on age groups within this dataset
- Most countries show a similar risk distribution:
 Low risk (~53-54%) with most common category,
 Medium risk (~35-36%) follows, High risk (~10-11%) is the least frequent in most countries. This suggests that thyroid cancer risk is generally low in most global regions
- India Shows a Unique Risk Pattern High-risk individuals make up 32.86% of the population (significantly higher than other countries)
- Japan (10.06%) and South Korea (10.50%) have the lowest high-risk proportions

Collaboration & Outcomes

Outcomes

Are you happy with the final product? Yes

What do you hope to achieve in the next development cycle?

Fully utilize the dataset to its full potential

What would you do differently if you could start again?

We identified the bias but didn't take any action on it, we left the dataset as it was because of time constraints and no experience with handling this situation.

Development Problems

Problems that arose during development?:

Git collaboration

In group conflicts and resolutions?

Did you find any of the behaviour related content useful? Teamwork, problem solving etc?

Yes, mindful collaboration and problem-solving, work division

Interactivity: Overall good

Summary

Overall group dynamic: Good, friendly, professional

Overall satisfaction: 9/10 (-1 for

git)

What we learned: github collaboration

Our experiences: We had many issues with git commands but we had good troubleshooting and good mindset for the project

Q&A