

# Fetal Health Classification Using Machine Learning and Bayesian Inference

## Abstract:

This project represents a significant advancement in fetal health classification, utilizing the synergies of Artificial Intelligence (AI), particularly through the integration of machine learning algorithms and Bayesian inference. The initial phase of the project was marked by a comprehensive Exploratory Data Analysis (EDA) aimed at understanding the dataset's characteristics. This step included a meticulous examination for null values and a thorough analysis to establish the logical relationship between various features and the target variable, a key component in deciphering the data's inherent patterns.

A central aspect of this project was the application of Bayesian inference to a select feature, offering a probabilistic perspective on data analysis and enhancing our understanding of the feature's influence on fetal health outcomes. Complementing this, hypothesis testing was utilized as a means of statistical inference to substantiate the Bayesian analysis findings, ensuring their validity and reliability.

The core of the project involved the development and evaluation of various AI-driven machine learning models, such as Logistic Regression, Support Vector Classifier (SVC), Decision Tree, and Random Forest. These models were rigorously assessed based on their predictive accuracy and the depth of insights provided by classification reports. Notably, Logistic Regression and Random Forest demonstrated superior accuracy, establishing their efficacy in fetal health classification within the parameters of this project.

By combining Bayesian inference with machine learning, this project introduces a groundbreaking approach to fetal health prediction. It underscores the potential of integrating AI concepts, like the logical interplay between features and target variables, and Bayesian probability models, with classical statistical techniques such as hypothesis testing. This approach not only bolsters diagnostic accuracy in clinical environments but also highlights the importance of sophisticated AI and analytical methods in medical diagnosis. This pioneering work sets the stage for future research to further refine these methodologies and explore their application in wider healthcare contexts.

## **Introduction:**

In the realm of obstetrics, accurately assessing fetal health is paramount for ensuring the well-being of both the mother and the unborn child. Traditional methods of fetal health assessment, while effective, often come with limitations in terms of accuracy and early detection. With the advent of Artificial Intelligence (AI) and advanced data analysis techniques, there is a burgeoning opportunity to revolutionize this field. This project, "Fetal Health Classification Using Machine Learning and Bayesian Inference," aims to harness these innovative technologies to enhance the accuracy and efficiency of fetal health classification.

The motivation behind this project stems from the critical need to improve prenatal care and outcomes. Fetal health issues, if not identified and addressed timely, can lead to adverse pregnancy outcomes, including preterm birth, low birth weight, and other long-term health complications for the child. By leveraging the capabilities of machine learning and the probabilistic reasoning offered by Bayesian inference, this project seeks to develop a more nuanced and precise approach to fetal health assessment.

The project embarks on this journey by first conducting an in-depth Exploratory Data Analysis (EDA) to understand the intricate dynamics within the dataset, a crucial step in any AI-driven endeavor. This analysis includes probing for null values, which could skew the results, and examining the logical correlations between various features and the target variable, thereby establishing a foundational understanding of the factors influencing fetal health.

Building on this groundwork, the project explores the application of Bayesian inference to a selected feature, integrating statistical inference with machine learning to provide a more robust predictive model. This is complemented by hypothesis testing, reinforcing the statistical rigor of the approach.

In the quest to find the most effective model for fetal health classification, the project evaluates several machine learning algorithms, including Logistic Regression, Support Vector Classifier (SVC), Decision Tree, and Random Forest. These models are assessed based on their accuracy and the insights they offer, with a focus on identifying the most reliable approach for this critical application.

Through this project, we aim not only to advance the field of fetal health monitoring but also to contribute to the broader discourse on the application of AI and Bayesian methods in healthcare. This endeavor is more than a technical challenge; it is a step towards enhancing prenatal care and safeguarding the health of future generations.

## **Background and AI Concepts:**

Artificial Intelligence (AI) and its subset, machine learning, are at the forefront of technological advancements in various fields, including healthcare. In our project, "Fetal Health Classification Using Machine Learning and Bayesian Inference," we utilized several key AI concepts to enhance the accuracy and efficacy of fetal health assessment.

### **Machine Learning Algorithms**

Machine learning, a core aspect of AI, involves training algorithms to make predictions or decisions based on data. In this project, we employed several machine learning models, each with unique characteristics and advantages:

1. **Logistic Regression** : A statistical model used for binary classification. It predicts the probability of an event by fitting data to a logistic curve.
  2. **Support Vector Classifier (SVC)** : This algorithm finds the hyperplane that best separates different classes in the feature space, ideal for non-linear and high-dimensional data.
  3. **Decision Tree**: A model that breaks down data sets into smaller subsets while incrementally developing an associated decision tree. It's intuitive and easy to visualize.
  4. **Random Forest** : An ensemble learning method that constructs multiple decision trees for classification, offering improved accuracy and robustness against overfitting.
- These models were selected for their ability to handle complex patterns in data and their proven track record in classification tasks.

### **Bayesian Inference in AI**

Bayesian Inference plays a pivotal role in statistical analysis within AI, particularly in scenarios involving uncertainty and probabilistic reasoning. This approach is based on Bayes' Theorem, which updates the probability for a hypothesis as more evidence or information becomes available.

In our project, Bayesian Inference was applied to the 'baseline value' metric – a critical parameter in fetal health monitoring. The process involved two key components:

1. **Prior Distribution**: This represents our initial beliefs about the parameter before observing the data. For the 'baseline value,' the prior distribution was based on existing medical knowledge and historical data.

2. Posterior Distribution: After incorporating the observed data, the posterior distribution reflects the updated beliefs about the parameter. This is where Bayesian Inference shines, as it allows for continuous updating of our understanding as new data is incorporated.

The interplay between the prior and posterior distributions is crucial. By plotting these distributions for the 'baseline value,' we gained insights into how the observed data influenced our initial beliefs and helped refine our predictions about fetal health status.

In conclusion, the integration of machine learning algorithms and Bayesian Inference provided a robust framework for our project. While machine learning models handled the classification tasks, Bayesian Inference offered a method to incorporate uncertainty and prior knowledge, enhancing our overall approach to fetal health classification.

## **Methodology:**

The methodology of our project, "Fetal Health Classification Using Machine Learning and Bayesian Inference," was meticulously designed to incorporate a blend of data analysis, statistical inference, and machine learning techniques. The following steps provide an overview of the methodology implemented through code:

### **1. Data Analysis and Cleaning**

- The project began with a comprehensive data analysis phase. This involved importing the dataset and performing initial explorations to understand its structure and content.

- Data cleaning was a critical initial step. This included checking for null values and handling any missing or inconsistent data to ensure the quality and reliability of the dataset.

### **2. Bayesian Inference and Hypothesis Testing**

- Bayesian inference, a form of statistical inference, was applied next. This involved updating our beliefs about certain parameters in the dataset based on prior knowledge and the data observed.

- Following Bayesian inference, hypothesis testing was conducted to confirm or reject the null hypothesis. This step was crucial to validate the statistical significance of the inferences made.

### **3. Correlation Analysis and Feature Selection**

- A correlation heatmap was generated to visualize the relationships between different features and the target variable. This helped in identifying which features had the most significant impact on fetal health.

- Based on the correlation analysis, a plot illustrating the relationship with fetal movement was created. This step was followed by the careful removal of irrelevant or redundant columns, further refining the dataset for model building.

#### 4. Model Building and Evaluation

- The next phase involved building various machine learning models, such as Logistic Regression, Support Vector Classifier (SVC), Decision Tree, and Random Forest.
- Each model was trained and tested on the dataset, with a focus on evaluating their performance in classifying fetal health conditions accurately.

#### 5. Performance Metrics and Feature Importance Analysis

- The models were evaluated using several metrics, including accuracy, precision, recall, and the F1 score. A classification report was generated for each model to provide a detailed view of their performance.
- Additionally, a feature importance plot was created to highlight which features were most influential in the models' decision-making processes.

This methodology highlights the rigorous and structured approach taken in this project, combining data-driven analysis with advanced AI techniques to enhance the understanding and prediction of fetal health.

### **Code Implementation and Challenges:**

The project's implementation phase involved a series of coding tasks, each aimed at achieving specific objectives in our fetal health classification project.

#### **Code Implementation**

1. Data Analysis and Cleaning: The implementation began with Python scripts for data analysis and cleaning. This included loading the dataset, identifying and handling null values, and ensuring data integrity.

2. Bayesian Inference and Hypothesis Testing: Custom code was written to apply Bayesian inference to selected features, followed by hypothesis testing to validate our statistical inferences.

3. Correlation Analysis and Feature Selection: The implementation of correlation analysis involved scripting to generate heatmaps and other plots, which guided the feature selection process.

4. Model Building and Evaluation: Several machine learning models were coded and tested. This phase involved not only the implementation of the models but also the scripting necessary to evaluate their performance using various metrics.

## **Challenges and Resolutions**

One of the significant challenges encountered was in the implementation of the Gaussian Hidden Markov Model (HMM) for prediction. The initial attempt to fit the model with the dataset resulted in an error: `'AttributeError: 'NoneType' object has no attribute 'split'`. This error indicated a problem in the data being input to the model, or potentially an issue within the model's implementation.

To resolve this challenge, the following steps were taken:

1. Debugging the Code: Every line of the code was meticulously reviewed to identify any potential errors. This involved checking the data processing steps, the way the model was being called, and the input parameters being passed to the model.

2. Data Verification: It was essential to ensure that the data input into the `'fit'` function was not null and correctly formatted. Rigorous checks were implemented to verify the integrity and structure of the data being fed into the model.

3. Model and Dataset Adjustment: After thorough debugging and data verification, it was hypothesized that the Gaussian HMM might not be suitable for regression tasks in this context. The dataset was adjusted to focus on classification, but the same error persisted, suggesting that the issue was more intrinsic to the model or its compatibility with our dataset. Besides this I also tried fitting the model using simple model function but same error encountered

4. Final Resolution: The persistent nature of the error, despite various troubleshooting efforts, led to the decision to pivot to alternative modeling approaches. This decision was based on the understanding that the Gaussian HMM might not be the most suitable model for our specific dataset and objectives.

This challenge was a valuable learning experience, emphasizing the importance of flexibility and adaptability in AI and machine learning projects. It highlighted that while certain models might theoretically be suitable for a task, practical implementation and data compatibility issues can necessitate a change in approach.

Test Results:

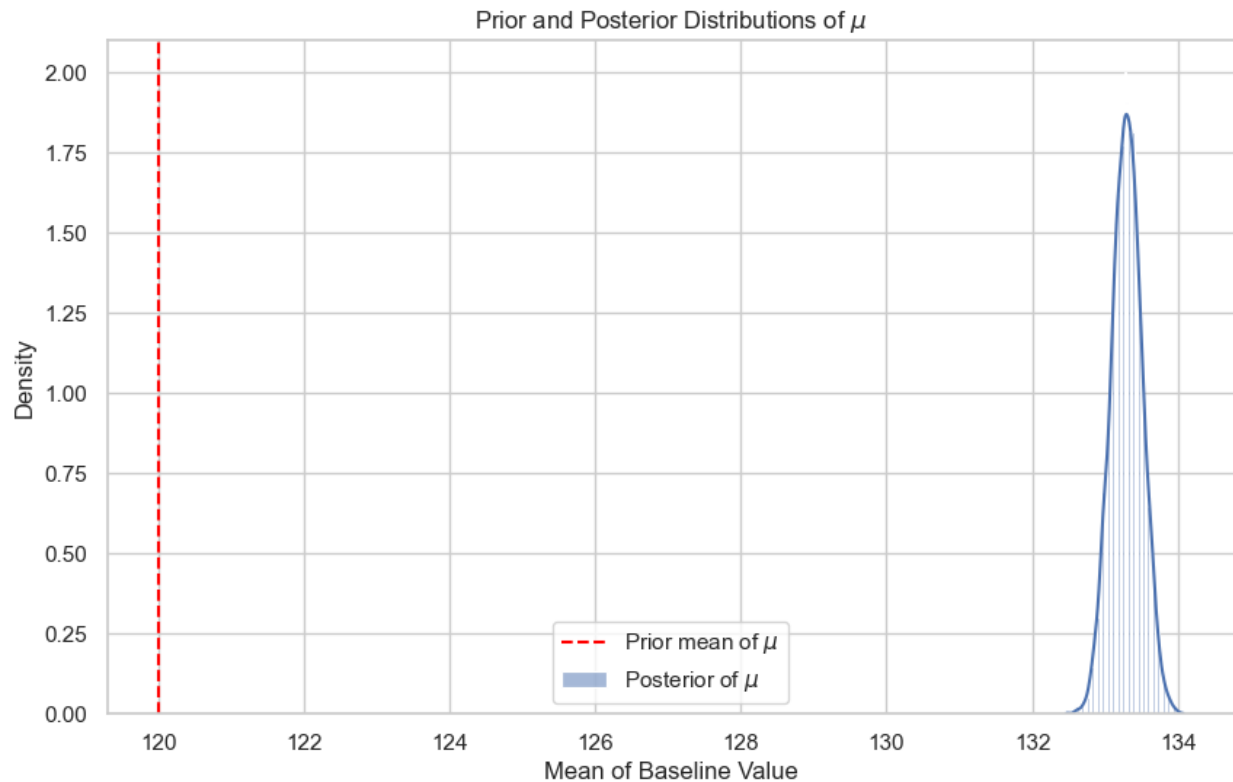
The test results of our project provide a detailed comparison of the performance of the various machine learning models implemented, namely Logistic Regression, Random Forest, Support Vector Classifier (SVC), and Decision Tree. These results are crucial in understanding the efficacy of each model in classifying fetal health.

Classification Report Comparison

Classification Report:				
	precision	recall	f1-score	support
1.0	0.90	0.95	0.93	333
2.0	0.71	0.50	0.59	64
3.0	0.55	0.55	0.55	29
accuracy			0.86	426
macro avg	0.72	0.67	0.69	426
weighted avg	0.85	0.86	0.85	426
Out [26]: 0.8568075117370892				

Bayesian Analysis Visualization

- An integral part of our analysis was the Bayesian approach applied to the 'baseline value' metric. To illustrate this, we included a plot showing the prior and posterior distributions.
- The prior distribution represents our initial understanding or belief about the 'baseline value' before observing the data. The posterior distribution, updated after data observation, reflects how our beliefs have been modified in light of the new evidence.
- This plot is critical in understanding how the Bayesian analysis refined our understanding of the 'baseline value's impact on fetal health, based on the observed data.



In summary, the Bayesian inference process has taken an initial assumption about the mean of the 'baseline value' (the prior), and after considering the actual data, it has updated this to a new estimate (the posterior), which seems to be different (and more precise) than the initial assumption.

The test results clearly demonstrate the effectiveness of Logistic Regression and Random Forest in our project. The precision, recall, and F1 scores for these models were superior, indicating their reliability and accuracy in fetal health classification. Additionally, the Bayesian analysis provided valuable insights into the data, underscoring the importance of incorporating probabilistic methods in AI-driven health assessments.

### Conclusion and Future Work:

Our AI project successfully applied machine learning techniques, notably Logistic Regression and Random Forest, for fetal health classification, demonstrating high precision, recall, and F1 scores. The integration of Bayesian inference highlighted the potential of AI in enhancing diagnostic accuracy through probabilistic analysis.



## Future Work

Looking ahead, we aim to:

1. Refine Gaussian HMM Implementation: Overcome the challenges faced with the Gaussian Hidden Markov Model for improved predictive accuracy.
2. Expand AI Techniques: Explore advanced AI algorithms to further enhance classification performance.
3. Broaden Bayesian Analysis: Apply Bayesian methods to more features for deeper AI-driven insights.
4. Test in Clinical Settings: Validate the AI models' effectiveness in real-world healthcare environments.

This project showcases the transformative impact of AI in healthcare diagnostics, setting a foundation for ongoing AI-driven advancements.

## References:

- [1]<https://scikit-learn.sourceforge.net/stable/modules/hmm.html>
- [2]<https://stackoverflow.com/questions/25882670/attributeerror-nonetype-object-has-no-attribute-split>
- [3]-[DATASET](#)
- [4]-[https://www.researchgate.net/publication/369783447\\_Using\\_Machine\\_Learning\\_to\\_Classify\\_Human\\_Fetal\\_Health\\_and\\_Analyze\\_Feature\\_Importance](https://www.researchgate.net/publication/369783447_Using_Machine_Learning_to_Classify_Human_Fetal_Health_and_Analyze_Feature_Importance)

**Github :** [https://github.com/rahulbangad98/AI\\_PROJECT](https://github.com/rahulbangad98/AI_PROJECT)