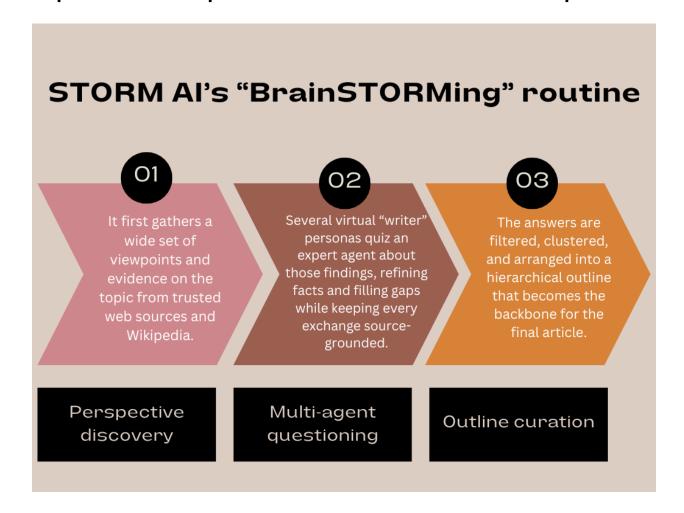
TOPIC

Al in predictive medical diagnostics to automatically diagnose possible health problems from all kinds of blood test panels



Brainstorming Process for this topic:

- 1. The Al tool first gathers information from a **Basic fact writer**, focusing on broadly covering the basic facts about the topic.
 - A. These fact-finding questions include the *main Al/ML algorithms and methods* currently being used in predictive medical diagnostics, particularly for analyzing blood test panels. Some ML algorithms, including Gaussian Naïve Bayes (GNB), k-nearest neighbors (KNN), support vector machines (SVM), decision trees (DT), and linear regression (LR), were seen to stand out. Supervised learning and Al algorithms can also analyze vast amounts of labeled clinical data, including historical medical records and laboratory results, to identify trends and predict future health risks

- B. The basic fact writer also looks for *current AI tools* being used to analyze lab results. The tools it found in the healthcare industry include <u>Lab-AI</u>, which utilizes Retrieval-Augmented Generation (RAG) to provide personalized normal ranges based on credible health data; deep neural network (DNN)-based automated recommendation system; <u>Docus AI Doctor</u>; <u>AI DiagMe</u>.
- C. The basic fact writer goes on to look for *primary challenges faced by healthcare providers when implementing AI systems*. It categorized the challenges and delved deep into each category: 1. Ethical Barriers, 2. Data Quality and Algorithmic Bias, 3. Regulatory Challenges, 4. Cognitive Errors and Dependency, 5. Workforce Challenges, 6. Social Implications.
- 2. Next, STORM AI created 3 personas with experience in different domains who could have an insight on this particular topic. They belong to different disciplines from where possible contributors could work on this topic:

A. Sarah Patel:

- An ethicist with a focus on health technology. She will address the ethical
 considerations and potential biases involved in using AI for medical diagnostics,
 as well as patient privacy concerns and the implications of relying on
 machine-generated diagnoses.
- Bias in training data can bake systemic inequalities into models, leading to unequal or inaccurate diagnoses for underrepresented groups.
- Limited dataset diversity pushes one size fits all predictions that miss important physiological differences and widen health disparities.
- Large, sensitive health datasets heighten privacy, consent, and transparency risks, calling for strict safeguards and clear accountability in Al use.

B. Dr. Emily Chen:

- A physician specializing in hematology who has experience using Al tools in clinical practice. She will focus on the clinical implications of Al in predictive diagnostics, discussing how these technologies can enhance patient care and accuracy in diagnosis.
- Key AI tools include classical machine learning models such as support vector machines, naive Bayes, decision trees, logistic regression, K nearest neighbors, plus deep neural networks like artificial and convolutional neural networks.
- These algorithms power predictive analytics that mine electronic medical records and feed automated decision support systems which suggest interpretations and optimal test selections.
- Some platforms also fuse laboratory panels with imaging data and combine supervised with unsupervised learning to reveal subtle patterns and raise diagnostic accuracy.

C. Mark Thompson:

- A data scientist working in healthcare analytics. He will delve into the technical side of Al algorithms, explaining how they process data from blood test panels and the importance of machine learning in identifying health trends.

- Support Vector Machines lead with strong classification in high dimensional data, often achieving accuracy above ninety seven percent for specific conditions.
- Artificial Neural Networks capture complex nonlinear relations in large panels, while interpretable Decision Trees and Logistic Regression offer clear rationale for binary or multiclass diagnoses.
- K Nearest Neighbors and Gaussian Naive Bayes suit preliminary screens on modest datasets, and ensembles like Random Forests enhance robustness by combining many tree models.

