NutriScopeAl V2 Workflow Chart

The workflow for NutriScopeAl V2 can be visualized as a sequential process with distinct modules handling specific tasks. The diagram below illustrates the flow from user interaction to the final report display.

1. Frontend Interaction

The process begins with the **frontend**, specifically frontend/pages/upload.py, where a user like Alice uploads her lab report PDF. This file, along with metadata (name, age, sex), is submitted to the backend through the /api/analyze endpoint. The frontend/pages/utils.py file ensures that API requests are properly authenticated. Once the analysis is complete, frontend/pages/report_details.py fetches and displays the final report, including a summary table and meal plan cards.

2. Backend API Orchestration

The **backend API**, managed by backend/app/api/routes.py, acts as the central hub. It receives the uploaded file from the frontend and orchestrates the entire analysis pipeline. This endpoint handles the initial request, triggering the series of subsequent steps, including ingestion, normalization, and AI summarization.

3. Ingestion & Parsing

This step is critical for reading the user's data. The backend/app/ingest/parser.py module extracts the essential information—test names, values, and units—from the PDF. If the PDF is a low-quality scan, the backend/app/ingest/extract.py provides an **OCR (Optical Character Recognition) fallback**, ensuring data can be extracted from image-based reports.

4. Normalization

After extraction, the raw data is cleaned and standardized. The backend/app/normalize/normalized_values.py maps extracted test names to a canonical form (e.g., "Hb" to "Hemoglobin"). Simultaneously, backend/app/normalize/unit_normalization.py standardizes all units, converting them to a consistent format (e.g., mg/dL to mmol/L), while backend/app/normalize/aliases.py handles various synonyms for test names.

5. Knowledge Base & Evaluation

With the data normalized, it's cross-referenced against a **knowledge base (KB)**. The backend/app/kb/loader.py module loads standard reference ranges for each lab test. The

backend/app/core/evaluator.py then flags each value as **Low**, **Normal**, or **High** by comparing the user's result to these reference ranges. This evaluation provides the foundational context for the AI summarization step.

6. Al Summarization & Meal Planning

This is the core of the NutriScopeAI. The backend/app/summarize/llm.py module uses the Groq LLM API to generate a personalized report. Using the flagged test results and the knowledge base context, it produces:

- An explanation of why a test is important.
- Potential **reasons** for high or low values.
- Associated risks if the condition is untreated.
- A personalized **meal plan** with food suggestions to address the flagged issues (e.g., iron-rich foods for low hemoglobin).

7. Storage & Persistence

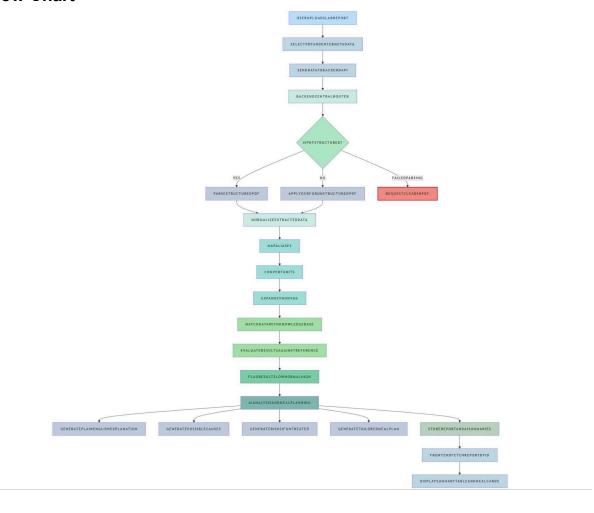
Once the AI-generated report is complete, the backend/app/storage/reports_store.py module saves the full report—including the original data and the AI-generated summaries—to a **JSON file** (backend/app/storage/reports.json). This ensures the data is durable and can be retrieved later by the user or the system.

8. Frontend Display

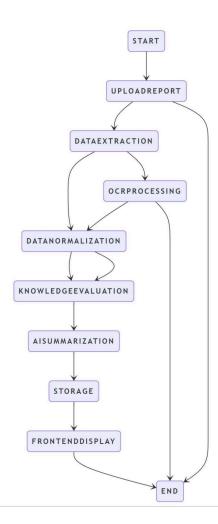
Finally, the completed report is ready to be displayed. The frontend fetches the stored report via the /api/report/{rid} endpoint. The frontend/pages/report_details.py file then formats and presents the information in a user-friendly manner, showing a structured summary table and a clear, easy-to-read meal plan.

This shows the flow from Frontend \rightarrow API \rightarrow Parsing \rightarrow Normalization \rightarrow KB \rightarrow Core Evaluation \rightarrow Al Summarization \rightarrow Storage \rightarrow Back to Frontend with test reports feeding into parsing.

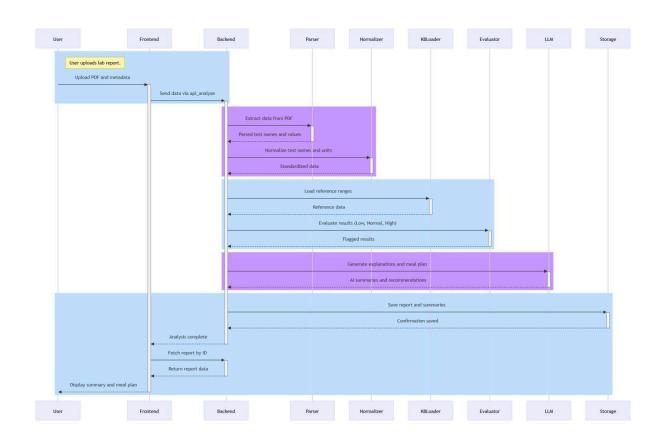
9. Flow Chart



10. State Diagram



11. Sequence Diagram



12. Class Diagram

