

Medical Report Simplifier

An AI-powered backend service that processes medical reports (text or images) and provides patient-friendly explanations using OCR, text normalization, and Google Gemini AI.

Features

- **Multi-format Input:** Supports both text input and image uploads (PNG, JPG, PDF)
- **OCR Processing:** Extracts text from medical report images with confidence scoring
- **Text Normalization:** Standardizes medical test names, values, units, and reference ranges
- **AI-Powered Explanations:** Uses Google Gemini to generate patient-friendly summaries
- **AI-Powered Validation:** Prevents hallucinated results using semantic comparison
- **4-Step Pipeline:** OCR → Normalization → AI Explanation → Final Output

Tech Stack

- **FastAPI** - Modern web framework for APIs
- **Google Gemini API** - AI for text processing and explanations
- **Tesseract OCR** - Text extraction from images
- **Pydantic** - Data validation and serialization
- **Python 3.9+** - Core programming language

Setup Instructions

Prerequisites

1. **Python 3.9+** installed
2. **Tesseract OCR** installed:

```
# Ubuntu/Debian
sudo apt-get install tesseract-ocr

# macOS
brew install tesseract

# Windows - Download from: https://github.com/UB-
Mannheim/tesseract/wiki
```

3. **Google Gemini API Key:**

- Go to [Google AI Studio](#)
- Create a new API key
- Copy the API key for configuration

Installation

1. Clone the repository:

```
git clone <your-repo-url>
cd medical-report-simplifier
```

2. Create virtual environment:

```
python -m venv venv
source venv/bin/activate # On Windows: venv\Scripts\activate
```

3. Install dependencies:

```
pip install -r requirements.txt
```

4. Configure environment:

```
cp .env.example .env
```

Edit `.env` and add your Google Gemini API key:

```
GEMINI_API_KEY=your_api_key_here
```

Running the Application

1. Start the server:

```
python3.9 -m uvicorn app.main:app --reload --port 8000
```

2. Access the API:

- API Documentation: <http://localhost:8000/docs>
- Health Check: <http://localhost:8000/health>

API Usage Examples

1. Process Text Input

```
curl -X POST "http://localhost:8000/process-text" \
  -H "Content-Type: application/json" \
  -d '{
    "text": "CBC: Hemoglobin 10.2 g/dL (Low), WBC 11,200 /uL (High)"
  }'
```

2. Process Image Input

```
curl -X POST "http://localhost:8000/process-image" \
  -H "Content-Type: multipart/form-data" \
  -F "file=@medical_report.png"
```

Expected Response Format

```
{
  "tests": [
    {
      "name": "Hemoglobin",
      "value": 10.2,
      "unit": "g/dL",
      "status": "low",
      "ref_range": {"low": 12.0, "high": 15.0}
    },
    {
      "name": "WBC",
      "value": 11200,
      "unit": "/uL",
      "status": "high",
      "ref_range": {"low": 4000, "high": 11000}
    }
  ],
  "summary": "Low hemoglobin and high white blood cell count.",
  "status": "ok"
}
```

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Architecture

```
app/
├─ main.py           # FastAPI application entry point
├─ models/
│   └─ schemas.py    # Pydantic models for request/response
├─ services/
│   └─ ocr_service.py # OCR and text extraction
│   └─ normalization_service.py # Medical test normalization
```

```
|   ├── ai_service.py      # Gemini AI integration
|   └── medical_data.py    # Medical reference data
└── api/
    ├── endpoints.py      # API route handlers
    └── core/
        ├── config.py     # Configuration management
        └── utils.py       # Utility functions
```

Testing

Run tests with:

```
pytest tests/ -v
```

Deployment

Local Demo with ngrok

1. **Install ngrok:** <https://ngrok.com/download>
2. **Start the FastAPI server:**

```
uvicorn app.main:app --host 0.0.0.0 --port 8000
```

3. **Expose with ngrok:**

```
ngrok http 8000
```

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Cloud Deployment

The application can be deployed to:

- **Railway:** Connect GitHub repo and deploy
- **Render:** Connect GitHub repo and deploy
- **Heroku:** Use Procfile for deployment

AI-Powered Hallucination Validation

The Problem

Traditional validation methods use simple string matching, which can miss:

- Fabricated test results that weren't in the original data
- Incorrect values or units
- Completely invented medical tests

Our Solution: Semantic Context Comparison

We use AI to compare the **semantic context** between original text and normalized results:

```
def validate_against_hallucination(self, original_tests: List[str],
                                   normalized_tests: List[NormalizedTest])
    -> Tuple[bool, str]:
    """
        Use AI to validate that normalized tests don't contain hallucinated
        information
        by comparing semantic context between original and normalized data
    """
```

How It Works

1. **Input Comparison:** AI compares original medical text with normalized results
2. **Semantic Analysis:** Detects context mismatches beyond simple string matching
3. **Confidence Scoring:** Uses confidence thresholds (>0.7) for reliability
4. **Medical Standardization:** Allows reasonable conversions (e.g., "Hgb" → "Hemoglobin")
5. **Fabrication Detection:** Identifies completely invented tests or wrong values

Example Scenarios

✅ **Valid:** Original: "Hgb 10.2 g/dL Low" → Normalized: "Hemoglobin: 10.2 g/dL (Low)"

❌ **Invalid:** Original: "Hgb 10.2 g/dL Low" → Normalized: "Cholesterol: 220 mg/dL (High)"

Testing the Validation

Run the demo script to see validation in action:

```
python demo_validation.py
```

Run validation-specific tests:

```
pytest tests/test_validation.py -v
```

Error Handling

The API includes comprehensive error handling:

- Invalid file formats
- OCR processing failures
- AI service errors
- Validation errors

- Hallucination detection with detailed error messages

Contributing

1. Fork the repository
2. Create a feature branch
3. Make your changes
4. Add tests
5. Submit a pull request

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