



# LungAI

Intelligent Lung Disease Detection System

## Comprehensive Project Documentation

|                   |   |
|-------------------|---|
| <b>Version</b>    | 1.0   |
| <b>Generated</b>  | 25 February 2026, 06:29 PM                        |
| <b>Project</b>    | Final Year AI Engineering Project                 |
| <b>Tech Stack</b> | Python · Flask · TensorFlow · MobileNetV2 · React |
| <b>Platform</b>   | Web Application (localhost:5001)                  |

■ For Educational & Engineering Demonstration Purposes Only

# 1. Project Overview

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**LungAI** is a Final Year AI Engineering Project that provides automated lung disease detection from chest X-ray images using deep learning. The system classifies X-ray images into four categories in real time and provides clinical insights, Grad-CAM explainability heatmaps, IoT-integrated vital signs, patient management, and downloadable PDF reports — all through a modern web interface.

## Key Capabilities:

- ✓ Automated 4-class chest X-ray classification (COVID-19, Pneumonia, Tuberculosis, Normal)
- ✓ Grad-CAM heatmap overlay for AI explainability (shows WHERE the model is looking)
- ✓ Severity scoring: Mild / Moderate / Severe based on confidence %
- ✓ Clinical risk badge: Low / Moderate / Severe / Critical Risk
- ✓ AI-generated medical insights, diet recommendations & medication guidance
- ✓ IoT-simulated real-time vitals: Heart Rate, SpO<sub>2</sub>, Temperature, Respiratory Rate
- ✓ PACS History Timeline — view all past analyses per patient
- ✓ Downloadable PDF clinical report per analysis
- ✓ Doctor Portal & Patient Portal with login/role-based access
- ✓ Google OAuth authentication support

# 2. Training Dataset

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The model was trained on a curated multi-class chest X-ray dataset organised into four disease categories. The dataset follows a standard train / val / test split stored under **data/chest\_xray\_multi/**.

## Dataset Structure:

| Split        | COVID-19      | Normal        | Pneumonia     | Tuberculosis | Total (approx.)      |
|--------------|---------------|---------------|---------------|--------------|----------------------|
| Train        | ~1,800        | ~1,800        | ~3,900        | ~700         | ~8,200 images        |
| Val          | ~200          | ~200          | ~400          | ~80          | ~880 images          |
| Test         | ~200          | ~234          | ~390          | ~80          | ~900 images          |
| <b>TOTAL</b> | <b>~2,200</b> | <b>~2,234</b> | <b>~4,690</b> | <b>~860</b>  | <b>~9,980 images</b> |

*Note: Exact counts depend on the dataset version used. Class imbalance (Pneumonia >> Tuberculosis) is handled automatically using computed class weights during training.*

## Data Augmentation Applied During Training:

- Rotation: ±10° — simulates patient positioning variation
- Width & Height Shift: ±10% — accounts for off-centre X-rays
- Shear Range: 0.1 — slight distortion tolerance
- Zoom Range: ±20% — scale invariance
- Brightness Range: [0.8 – 1.2] — exposure variation

- Horizontal Flip: True — bilateral symmetry of lungs
- Fill Mode: nearest — border fill after transformation

All images are resized to **224 × 224 pixels** and normalised using MobileNetV2's preprocessing function (scales pixel values to the range **[-1, 1]**).

### 3. Algorithms & Model Architecture

#### 3.1 Primary Algorithm — Transfer Learning with MobileNetV2

The core classification algorithm is **Transfer Learning** using **MobileNetV2** (pre-trained on ImageNet). MobileNetV2 was chosen for its balance of accuracy and computational efficiency — making it deployable on standard hardware without a GPU.

| Layer / Component      | Details   |
|------------------------|---|
| Base Model             | MobileNetV2 (ImageNet weights, include_top=False)   |
| Input Shape            | 224 × 224 × 3 (RGB)                                 |
| GlobalAveragePooling2D | Reduces spatial feature maps to 1D vector           |
| Dense (512 units)      | Activation: ReLU — learns disease-specific features |
| BatchNormalization     | Normalises activations for stable training          |
| Dropout (0.5)          | 50% neuron drop — prevents overfitting              |
| Dense (128 units)      | Activation: ReLU — further feature compression      |
| Dropout (0.3)          | 30% neuron drop — additional regularisation         |
| Dense (4 units)        | Activation: Softmax — probability per class         |
| Output Classes         | COVID19   NORMAL   PNEUMONIA   TUBERCULOSIS         |

#### 3.2 Two-Phase Training Strategy

| Phase                        | Epochs | Learning Rate   | What is trained                      | Optimizer |
|------------------------------|--------|---|--------------------------------------|-----------|
| Phase 1 — Feature Extraction | 10     | 0.001   | Only top custom layers (base frozen) | Adam      |
| Phase 2 — Fine-tuning        | 20     | Top 100 custom layers + last layers of MobileNetV2 (first 100 base layers frozen) | Adam                                 | Adam      |

#### 3.3 Other Algorithms & Techniques Used

| Algorithm / Technique          | Purpose  | Library               |
|--------------------------------|--|-----------------------|
| Softmax Classification         | Converts logits to probability distribution (4 classes)      | Keras                 |
| Categorical Cross-Entropy Loss | Multi-class training loss function                           | Keras                 |
| Adam Optimiser                 | Adaptive gradient descent for weight updates                 | TensorFlow            |
| Class Weight Balancing         | Corrects for dataset imbalance (Pneumonia >> TB)             | scikit-learn / custom |
| Early Stopping                 | Stops training when val_loss stops improving (patient Keras) | Keras                 |
| ReduceLROnPlateau              | Reduces LR by 0.2x when val_loss plateaus (patient Keras)    | Keras                 |
| ModelCheckpoint                | Saves best model weights by val_accuracy                     | Keras                 |

|  |  |                     |
|--|--|---------------------|
| Grad-CAM (Gradient-weighted Class Activation Maps) | Generates visual heatmaps showing which lung regions influenced the prediction | TensorFlow + OpenCV |
| Batch Normalisation                                | Normalises layer activations — faster, stable training                         | Keras               |
| Dropout Regularisation                             | Randomly disables neurons to prevent overfitting                               | Keras               |
| MobileNetV2 Preprocessing                          | Scales pixel values to [-1, 1] for ImageNet weights                            | Keras               |
| GradientTape                                       | Tracks gradients for Grad-CAM computation                                      | TensorFlow          |
| OpenCV COLORMAP_JET                                | Applies colour heatmap to the Grad-CAM output                                  | OpenCV (cv2)        |

## 4. How the Project Works

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### End-to-End Prediction Pipeline:

#### Step 1 — Upload

Doctor/user registers or selects a patient, then uploads a chest X-ray image (PNG/JPG/JPEG, max 16 MB) through the web interface at localhost:5001.

#### Step 2 — File Validation & Storage

Flask validates file type and saves it securely with a UUID filename to static/uploads/. The file path is stored relative to the static folder.

#### Step 3 — CNN Inference

The image is resized to 224x224 pixels, preprocessed (pixel values scaled to [-1,1]), and passed through the MobileNetV2-based model (lung\_model\_multi.h5). Outputs: a 4-element softmax probability vector → argmax gives the predicted class.

#### Step 4 — Grad-CAM Generation

Using TensorFlow GradientTape, gradients of the predicted class score are computed with respect to the last Conv2D layer's output. These are averaged (Global Average Pooling) and used to weight the feature maps, producing a heatmap that is overlaid on the X-ray using OpenCV.

#### Step 5 — Medical Insights

Based on the predicted class and confidence, the system looks up AI insights, clinical description, diet plan, and medication recommendations from medical\_insights.py.

#### Step 6 — Severity & Risk

Severity is computed from confidence:  $\geq 85\%$  → Severe,  $\geq 65\%$  → Moderate,  $< 65\%$  → Mild. Clinical risk is derived from the combination of severity and prediction class.

#### Step 7 — Database Storage

Patient info and the XRay report (prediction, confidence, severity, image path, heatmap path) are stored in the SQLite database via Flask-SQLAlchemy.

#### Step 8 — Result Display

The result.html template renders: AI prediction badge, confidence bar, Grad-CAM viewer, probability chart, medical insights, diet & medication cards, IoT vitals, and PACS history.

#### Step 9 — PDF Export

The /download-report/ route calls report\_generator.py using ReportLab to generate a professional PDF clinical report and sends it as a file download.

### IoT Vitals Monitoring:

The system includes an IoT integration layer (iot/api.py) that fetches real-time simulated patient vitals (Heart Rate, SpO<sub>2</sub>, Temperature, Respiratory Rate) from /api/simulated-vitals/ every 3 seconds via JavaScript polling. Vitals are stored in the Vitals database table and displayed live with a clinical clock on

the result page.

## 5. Technology Stack

| Layer            | Technology                 | Version      | Purpose                        |
|------------------|----------------------------|--------------|--------------------------------|
| Web Framework    | Flask                      | 2.3.3        | HTTP routing, templating, API  |
| Deep Learning    | TensorFlow / Keras         | 2.13 / 2.13  | CNN training and inference     |
| Base Model       | MobileNetV2                | ImageNet     | Transfer learning backbone     |
| Image Processing | OpenCV (cv2)               | 4.8.1        | Grad-CAM overlay, image resize |
| Image Processing | Pillow                     | 10.0.1       | Image loading and conversion   |
| Data Science     | NumPy                      | 1.24.3       | Array operations               |
| ML Utilities     | scikit-learn               | 1.3.2        | Class weights, evaluation      |
| Visualisation    | Matplotlib                 | 3.7.3        | Training plots                 |
| Database ORM     | Flask-SQLAlchemy           | 3.0.5        | Database models & queries      |
| DB Migrations    | Flask-Migrate              | 4.0.5        | Schema version control         |
| Database         | SQLite (development)       | Built-in     | Patient & report storage       |
| Database         | PostgreSQL (production)    | via psycopg2 | Production DB (Supabase)       |
| Frontend         | React (TypeScript)         | Vite build   | Doctor & Patient portals       |
| Frontend         | HTML5 + Bootstrap 5        | 5.3.2        | Prediction result pages        |
| Auth             | Flask-Login + Google OAuth | —            | Role-based access control      |
| PDF Generation   | ReportLab                  | 4.0.4        | Clinical PDF report export     |
| Server           | Gunicorn                   | 21.2.0       | Production WSGI server         |
| Cors             | Flask-CORS                 | 4.0.0        | Cross-origin resource sharing  |

## 6. How to Run the Project (Terminal Guide)

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### Prerequisites:

- Python 3.9+ (check: python3 --version)
- Node.js 18+ (check: node --version)
- pip (Python package manager)
- Git (optional, for cloning)
- 4 GB+ RAM recommended for model loading

### Step 1 — Navigate to the project folder

```
cd "/Users/britto/Documents/Lung Disease Project/lung-disease-ai"
```

### Step 2 — Create and activate a virtual environment

```
python3 -m venv venv  
source venv/bin/activate # macOS / Linux  
venv\Scripts\activate # Windows
```

### Step 3 — Install Python dependencies

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

### Step 4 — Set up environment variables (copy .env.example to .env and fill in values)

```
cp .env.example .env  
# Then edit .env and set:  
# SECRET_KEY=your-secret-key  
# DATABASE_URL=sqlite:///lung_disease.db  
# GOOGLE_CLIENT_ID=... (optional)
```

### Step 5 — Initialise the database

```
flask db upgrade  
# OR for first-time setup without migrations:  
python -c "from app import app; from database.models import db;  
app.app_context().push(); db.create_all()"
```

### Step 6 — (Optional) Train the model

```
python model/train_multi_class.py  
# Place your dataset in: data/chest_xray_multi/train|val|test//
```

### Step 7 — Install frontend dependencies

```
cd frontend  
npm install  
cd ..
```

## Step 8 — Run the full project (Backend + Frontend together)

```
npm run dev
```

# This runs both Flask (port 5001) and React dev server concurrently

## Step 8 (Alternative) — Run backend only

```
python app.py
```

# Flask runs at: http://localhost:5001

## Step 9 — Open in browser

```
http://localhost:5001
```

# Doctor Portal: http://localhost:5001/doctor

# Patient Portal: http://localhost:5001/patient/

## Useful Commands:

| Task                           | Command   |
|--------------------------------|---|
| Generate a PDF report via CLI  | python generate_docs_pdf.py                           |
| Run IoT vitals simulator       | python sim_iot.py                                     |
| Inspect trained model layers   | python model/list_layers.py                           |
| Export model info to JSON      | python model/export_model_info.py                     |
| Apply a new DB migration       | flask db migrate -m 'description' && flask db upgrade |
| Run with Gunicorn (production) | gunicorn -w 4 -b 0.0.0.0:5001 'app:app'               |

## 7. Project Folder Structure

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```
lung-disease-ai/
  app.py ← Main Flask application & all routes
  config.py ← Configuration (dev / prod / test)
  requirements.txt ← Python dependencies
  package.json ← Node scripts (npm run dev)
  sim_iot.py ← IoT vitals simulator
  model/
    lung_model_multi.h5 ← Trained CNN model (generated after training)
    train_multi_class.py ← Model training script (MobileNetV2)
    predict.py ← Inference + Grad-CAM generation
    medical_insights.py ← Disease insights, diet, medication data
    report_generator.py ← PDF clinical report generator (ReportLab)
    inspect_model.py ← Model layer inspection utility
  database/
    models.py ← SQLAlchemy models (Patient, XRayReport, Vitals, User)
    enable_rls.sql ← Row-Level Security SQL (for Supabase)
  iot/
    api.py ← IoT Blueprint: simulated vitals API
    templates/ ← Jinja2 HTML templates
    index.html ← Home / Upload page
    result.html ← Analysis result page
    login.html ← Login page
    register.html ← Register page
    product.html / technology.html
    static/ ← CSS/Style.css ← Custom styles
    js/main.js ← Frontend JS
    uploads/ ← Saved X-ray uploads
    heatmaps/ ← Grad-CAM heatmap images
    reports/ ← Generated PDF reports
    fe_assets/ ← React build output (Doctor/Patient portals)
    frontend/ ← React + TypeScript source (Vite)
    src/pages/ ← Dashboard, PatientPortal, DoctorPortal
    src/components/ ← Reusable UI components
    data/
      chest_xray_multi/ ← Training dataset (train/val/test//)
      migrations/ ← Flask-Migrate database migrations
```

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**Disclaimer:** LungAI is an engineering and educational demonstration project. All AI predictions are for informational purposes only and must NOT be used as a substitute for professional medical diagnosis. Always consult a qualified medical professional. Medication and diet information provided is general and informational — not a prescription.