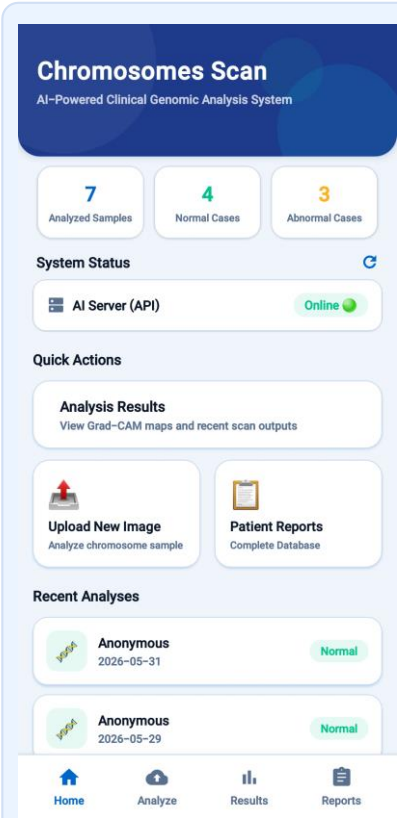


System Overview



Fig. 1 — Home Dashboard: system status, statistics, and quick-access panel.

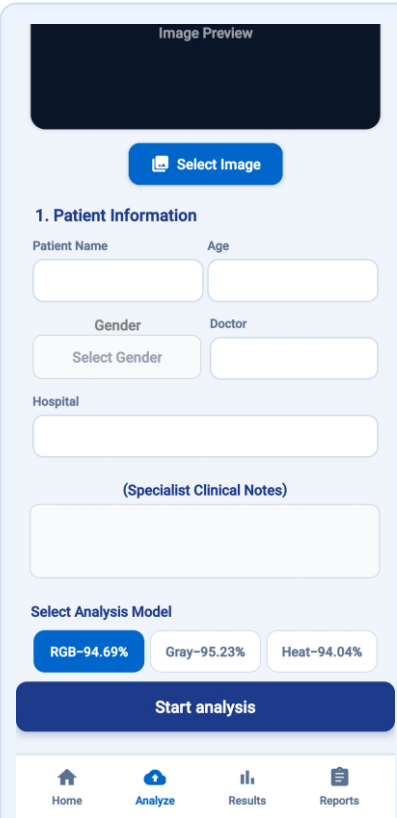
Step-by-Step Operating Procedure



01

Verify Server Synchronization

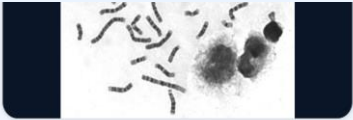
Upon initial boot, examine the top configuration panel. The application must reflect an **Online** connection status badge, indicating safe client-to-server HTTP synchronization.



02

Register Patient Biological Profile

Navigate to the **Analyze** navigation lane. Input dynamic validation demographics including Patient Name, Age, Supervising Clinician, and Hospital Center fields.



Select Image

1. Patient Information

Patient Name

عبدالرحمن

Age

40

Gender

Male

Doctor

عبدالله

Hospital

مستشفى X

(Specialist Clinical Notes)

Select Analysis Model

RGB-94.69%

Gray-95.23%

Heat-94.04%

Chromosomal testing is in progress...

Home

Analyze

Results

Reports

03

Acquire Chromosomal Image

1. Acquire Chromosomal Image

Tap Select Image to access your local device gallery and upload a high-resolution chromosomal spreadsheet (karyogram) acquired directly from your laboratory microscope setup.

2. Use Cytogenetic Rationale : Metaphase Chromosomes

In cytogenetic diagnostics, the captured chromosome images must be processed exclusively during the Metaphase stage of cell division due to the following critical biological factors:

Maximum Condensation: Chromosomes reach their peak coiling phase during this stage, becoming short, thick, and highly visible under digital microscopy.

Perfect Equatorial Alignment: The chromosomes line up uniformly along the cell's equatorial plate. This optimal spatial distribution prevents overlapping and crowding, which drastically improves AI segmentation boundaries and detection accuracy.

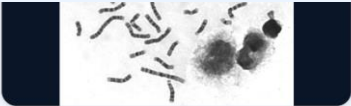
04

Select Target Neural Network

Choose the specialized CNN processing pipeline based on image contrast: **Gray-95.23%** for standard grayscale banding sheets, **RGB-94.69%** for color sheets, or **Heat-94.04%** mapping filters.

Chromosomes Scan — AI-Powered Clinical Genomic Analysis System

Page 3



Select Image

1. Patient Information

Patient Name

عبدالرحمن

Age

40

Gender

Male

Doctor

عبدالله

Hospital

مستشفى X

(Specialist Clinical Notes)

Select Analysis Model

RGB-94.69%

Gray-95.23%

Heat-94.04%

Chromosomal testing is in progress...

Home

Analyze

Results

Reports

05

Invoke Automated Inference

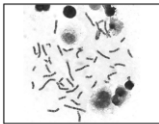
Invoke Automated Inference

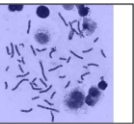
Trigger the Start Analysis command. The image is resized to 224×224 pixels and securely transmitted through secure channels for instant tensor evaluation.

1. Patient Information

Patient:	عبدالرحمن	Age:	40
Gender:	Male	Hospital:	مستشفى X
Doctor:	عبدالله	Date:	2026-05-31

2. AI Detection





(Original Scan)

(grade cam)

3. Analysis Findings & DNA/RNA Metrics

Status:	NORMAL
Total Chromosomes:	46
Abnormal Chromosome:	None
DNA Metrics:	1. Total Length: N/A 2. GC Content: N/A% 3. AT Content: N/A% 4. AT/GC Ratio: N/A
RNA Metrics:	1. Total Genes: 1524 2. Total RNA Transcripts: 3808 3. Average Gene Length: 34796.16
Gene Impact:	● No significant genomic variation detected. ● Normal karyotype profile mapped successfully.

Home

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06

Evaluate Clinical Insights

The system will redirect you to the **Results** display panel to evaluate classification markers, localization of anomalies, dynamic molecular statistics, and explainable Grad-CAM heatmaps.

07

Export Standard PDF Document

Tap **Download PDF Report** to activate the device printing spooler. This compiles a clean, multi-line, verified medical report with no layout clipping or formatting bugs.

08

History

	عبدالرحمن 2026-05-31	Normal
	Anonymous 2026-05-31	Normal
	Anonymous 2026-05-29	Normal
	Anonymous 2026-05-29	Abnormal
	Anonymous 2026-05-29	Normal
	Anonymous 2026-05-29	Abnormal
	Anonymous 2026-05-29	Normal
	Anonymous 2026-05-29	Abnormal

Persistent Database Caching

Close the interface or cycle system power safely. The non-volatile storage caching tier stores all patient logs under the **Reports (History)** tab, automatically modifying structural home panels.

			
Home	Analyze	Results	Reports

Grad-CAM Visual Explainability

Gradient-weighted Class Activation Mapping (Grad-CAM) provides spatial localization of the model's decision regions directly overlaid on the chromosomal scan. Warm regions (red/yellow) indicate high activation zones associated with the classified anomaly, enabling clinicians to visually validate the AI's reasoning.

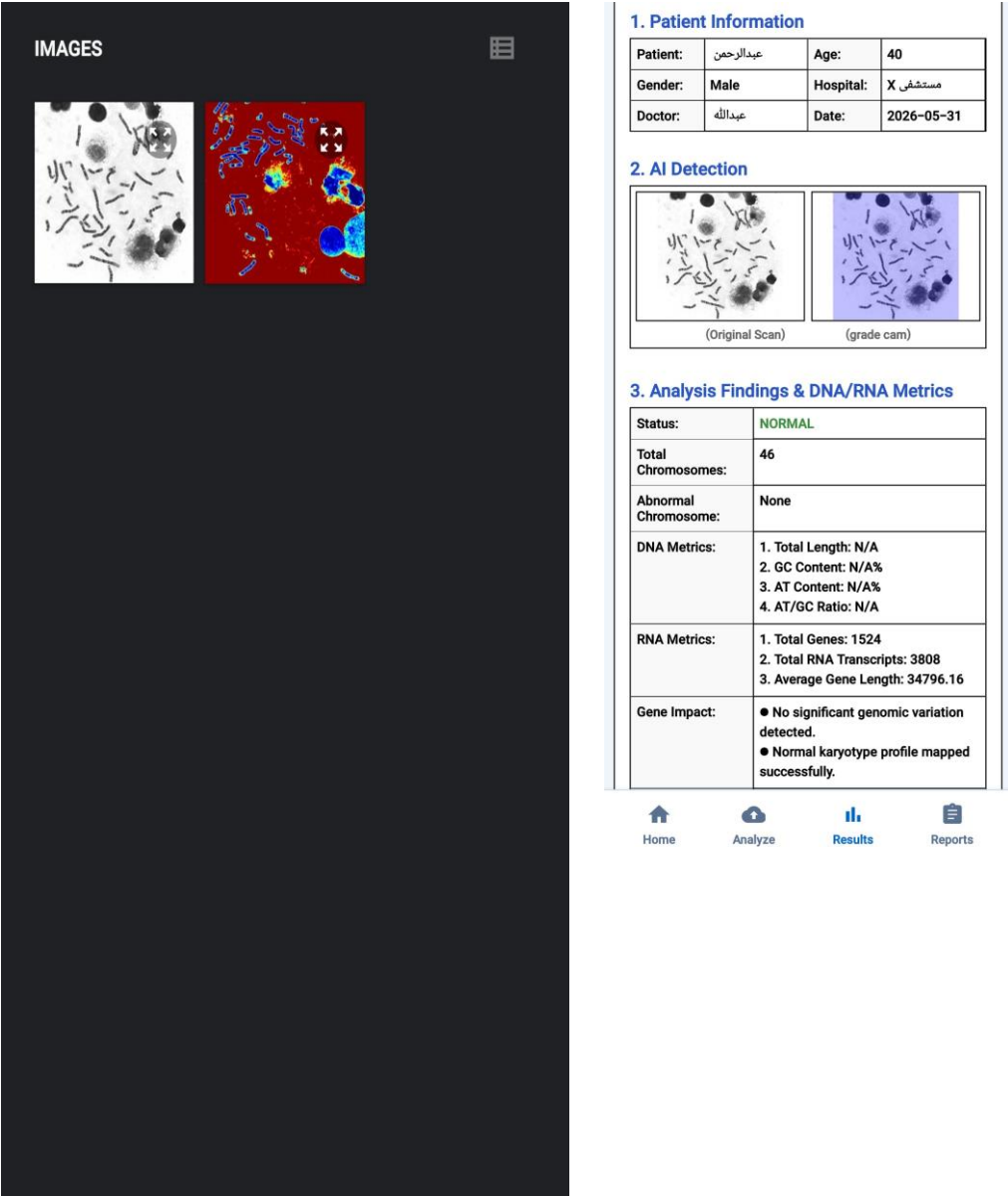


Fig. 2 — Left: Grad-CAM heatmap overlay on chromosomal scan. Right: Integrated results panel with molecular metrics and AI findings.

Technical Specifications

Parameter	Specification
Input Resolution	224 × 224 pixels (auto down-sampled)
Supported Formats	JPEG, PNG, BMP, TIFF (karyogram images)

CNN Architectures	RGB Model (94.69%), Grayscale (95.23%), Heatmap (94.04%)
Explainability Method	Gradient-weighted Class Activation Mapping (Grad-CAM)
Communication Protocol	HTTPS REST API — JSON response format
Local Storage	Non-volatile SQLite-based caching (AsyncStorage)
Report Output	Structured PDF — Patient + AI findings + Grad-CAM
Normal Reference	46 chromosomes (diploid human karyotype)
Platform	Android 10+ (API Level 29)

Clinical Disclaimer: Chromosomes Scan is designed as a decision-support tool for trained cytogenetic professionals. All AI-generated findings must be validated by a licensed medical specialist before clinical use. Copyright © 2026 Chromosomes Scan — All Rights Reserved.