



Cairo University

جامعة القاهرة

Faculty of Computers & artificial intelligence

كلية الحاسبات والذكاء الاصطناعي

Bioinformatics Program

المعلومات الحيوية

Early detection of colorectal cancer using machine learning & deep learning

This documentation submitted as required for the degree of bachelors in Computers and artificial intelligence.

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Abstract

Colorectal cancer (CRC) begins in the colon or the rectum. These diseases can likewise be called colon malignant growth or rectal cancer, contingent upon where they start. Colon malignant growth and rectal disease are regularly assembled in light of the fact that they share many highlights practically speaking. Colorectal disease is second reason for death in many communities additionally early conclusion have a more prominent possibility of survival. So colonoscopy represents a very important diagnostic modality for screening for colorectal cancer, we address a current issue in medical picture handling, the discovery of colorectal disease from colonoscopy videos. As per overall malignant growth measurements, colorectal disease is perhaps the most well-known disease. The most common way of screening and the expulsion of pre-malignant cells from the digestive organ is an essential undertaking to date. The traditional manual process is dependent on the expertise of the medical practitioner. We have Two phases in this project The first stage we utilized SVM model with an accuracy of 87 % in second stage we utilized deep learning CNN models with some accuracies.so, The main objective of this project is early detection of colon cancer based on medical image processing extracted from colonoscopy videos.

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List of Abbreviations

Word	Abbreviation
Colorectal cancer	CRC
Support Vector machine	SVM
Convolutional Neural Networks	CNN
Hypertext Markup Language	HTML
Cascading Style sheet	CSS
Java Script	JS
<u>P</u> HP <u>H</u> ypertext <u>P</u> reprocessor	PHP

Chapter One
Introduction

Introduction

- **What is Cancer?**

The frame consists of about 15 trillion cells, and each day billions of cells wear out or are destroyed. Each time a cell is destroyed the body makes a brand new cell to exchange it, trying to create a cell that's an ideal copy of the cell that was destroyed because the replacement cell must be capable of performing the identical function because the destroyed cell.

During the complex process of replacing cells, many errors occur.

The chassis correct most mistakes, sometimes mistakes are made and not corrected, many of the uncorrected mistakes have little

effect on health.

If these mistakes allow the newly made cell to divide independent of the checks and balance that control normal cell growth, that cell can begin to multiply in an uncontrolled manner, when this happens, a tumor can develop.

- **What is colorectal cancer?**

Colorectal cancer is the cancer of colon and rectum. It starts when the process of the normal replacement colon lining cells deviate. Health care professionals are certain that colorectal cancer is not contagious (a person cannot catch the disease from a cancer patient). Some people are more likely to develop colorectal cancer than others.

- **What are the risk factors that increase the chance of colorectal cancer?**

1. Age :

Increasing age is the main risk factor for colorectal cancer. Around 90% of colorectal cancers are diagnosed after age 50.

2. Race:

African Americans have a higher incidence of colorectal cancer than people of other races.

3. Diet and colorectal cancer:

Diets high in fat have been shown in numerous research studies to predispose people to colorectal cancer.

In countries with high colorectal cancer rates, the fat intake by the population is much higher than in countries with low cancer rates. It is believed that the digestion of fat that occurs in the small intestine and the colon leads to the formation of cancer-causing chemicals (carcinogens). Likewise, research studies also reveal that diets high in vegetables and high-fiber foods such as whole-grain breads and cereals contain less fat that produces these carcinogens and may counter the effects of the carcinogens. Both effects would help reduce the risk of cancer.

4. Inherited syndromes:

Genetic syndromes passed through generations of your family can increase your risk of colon cancer.

5. A sedentary lifestyle:

If you're inactive, you're more likely to develop colon cancer. Getting regular physical activity may reduce your risk of colon cancer.

6. Diabetes:

People with diabetes and insulin resistance have an increased risk of colon cancer.

7. Obesity:

People who are obese have an increased risk of colon cancer and an increased risk of dying of colon cancer when compared with people considered normal weight.

8. Smoking:

People who smoke may have an increased risk of colon cancer.

9. Alcohol:

Heavy use of alcohol increases your risk of colon cancer.

10. Radiation therapy for cancer:

Radiation therapy directed at the abdomen to treat previous cancers increases the risk of colon and rectal cancer.

• What are the symptoms of Colorectal Cancer?

Symptoms of colorectal cancer are numerous and nonspecific.
They include:

- Fatigue,
- Weakness,
- Shortness of breath,
- Change in bowel habits,
- Narrow stools,
- Diarrhea or constipation,
- Red or dark blood in stool,
- Weight loss,
- Abdominal pain,
- Cramps or bloating.

- **What are the stages of Colon Cancer?**

- **Stage I:**

The cancer has grown through the superficial lining (mucosa) of the colon or rectum but hasn't spread beyond the colon wall or rectum.

- **Stage II:**

The cancer has grown into or through the wall of the colon or rectum but hasn't spread to nearby lymph nodes.

- **Stage III:**

The cancer has invaded nearby lymph nodes but isn't affecting other parts of your body yet.

- **Stage IV:**

The cancer has spread to distant sites, such as other organs for instance, to your liver or lung.

- **What is Machine Learning?**

is a field of understanding and building methods that 'learn', that methods that leverage data to improve performance on some set of tasks. It is seen as a part of artificial intelligence.

Machine Learning Algorithm:

algorithms build a model based on sample data, known as training data, in order to make predictions or decisions without being

It is used in a wide variety of applications, such as in medicine, email filtering, speech recognition, and computer vision, where it is difficult or unfeasible to develop conventional algorithms to perform the needed tasks.



TYPES OF MACHINE LEARNING

Machine Learning

- SUPERVISED LEARNING**
(TASK-DRIVEN)
- UNSUPERVISED LEARNING**
(DATA-DRIVEN)
- REINFORCEMENT LEARNING**
(LEARN FROM ERRORS)

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The main difference between the two types is that supervised learning is done using a ground truth, or in other words, we have

Fig 1.2 types of ML

Prior knowledge of what the output values for our samples should be. Therefore, the goal of supervised learning is to learn a function that, given a sample of data and desired outputs, best approximates

the relationship between input and output observable in the data.

Unsupervised learning, on the other hand, does not have labeled outputs, so its goal is to infer the natural structure present within a set of data point

Machine Learning can help in:

Prediction of cancer susceptibility – risk assessment prior to occurrence.

Prediction of cancer recurrence – likelihood of redeveloping.

Prediction of cancer survivability – life expectancy,

survival, progression, tumor-drug sensitivity.

Supervised Learning:

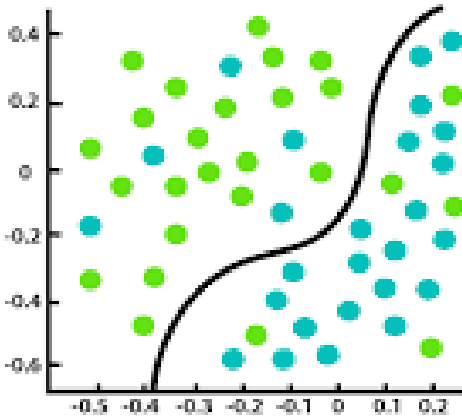
we are given a data set and already know what our correct output should look like, having the idea that there is a relationship between the input and the output.

Supervised learning problems are categorized into "regression" and "classification" problems.

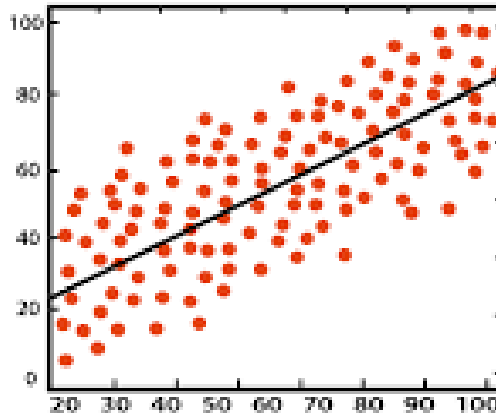
In a regression problem, we are trying to predict results within a continuous output, meaning that we are trying to map input variables to some continuous function.

In a classification problem, we are instead trying to predict results in a discrete output.

In other words, we are trying to map input variables into discrete categories



Classification



Regression

Fig1.3 Supervised Learning

Unsupervised Learning:

Unsupervised learning, on the other hand, allows us to approach problems with little or no idea what our results should look like.

We can derive structure from data where we don't necessarily know the effect of the variables. We can derive this structure by clustering the data based on relationships among the variables in the data.

With unsupervised learning there is no feedback based on the prediction results, i.e., there is no teacher to correct you

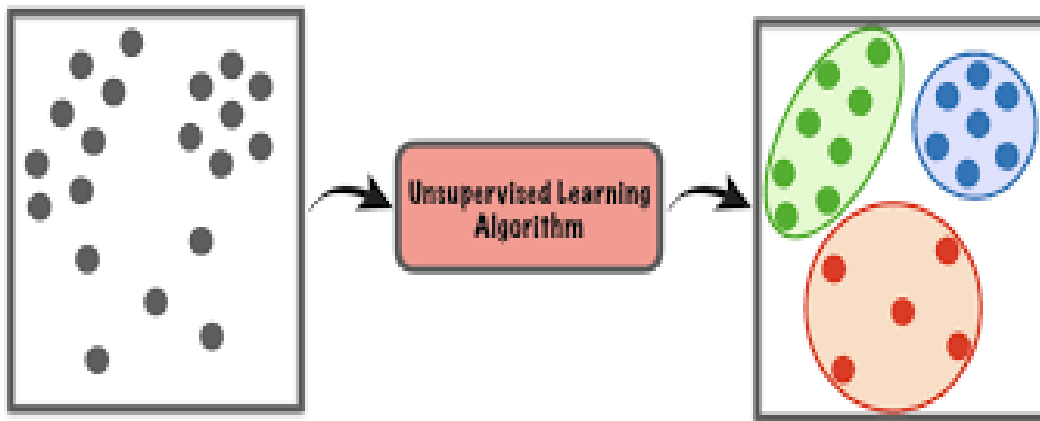


Fig 1.4

1.1 Motivation

Colorectal Cancer (CRC) is among the three most common cancers with more than 1.2 million new cases and about 600,000 deaths per year worldwide. The diagnosis of colorectal cancer at late stage makes the treatment more difficult and decreases to less than 10% chance of survival, on the contrary, early diagnosis of CRC is easier to treat and patient has 90% chance of survival. Using the medical pictures of various areas in the colon

Machine learning and deep learning models that predict whether the given picture is positive(has cancer) or negative(has no cancer)

1.2 Problem Definition

Currently, all detection of colorectal cancer based on traditional ways which are chemical and experimental studies as if your signs and symptoms indicate that you could have colon cancer, your doctor may recommend one or more tests and procedures, including:

Using a scope to examine the inside of your colon:

Colonoscopy uses a long, flexible and slender tube attached to a video camera and monitor to view your entire colon and rectum. If any suspicious areas are found, your doctor can pass surgical tools through the tube to take tissue samples for analysis and remove polyps.

Blood tests:

No blood test can tell you if you have colon cancer. But your doctor may test your blood for clues about your overall health, such as kidney and liver function tests. So, they detect cancer at late stage which decreases the probability of survival

1.3 Project Objective

The main objective of this project is early detection of colon cancer based on colonoscopy

Using a machine learning approach (SVM) to implement a classifier that can be trained on the labeled data, then it can classify the images into three different colonic locations and predict the case of that area whether it has a cancer or not

Using a deep learning approach (CNN) to implement model that can be trained on labeled data , so it can classify the images into three different colonic locations and predict the case of that area whether it has a cancer or not







1.4 Gantt chart of project time plan



Fig1.5 Gantt Chart

1.5 Project development methodology

Our project is about early detection of colorectal cancer first by machine learning specifically SVM model, second by deep learning and we used some models with different layers to give different accuracy. we present Kvasir, a dataset containing images from inside the digest, Kvasir is important for research on both single- and multi-disease computer aided detection. By providing it, we invite and enable multimedia researcher into the medical domain of detection and retrieval. The Kvasir dataset consists of images, annotated and verified by medical doctors (experienced endoscopists), including several classes showing anatomical landmarks, pathological findings or endoscopic procedures in the digest, hundreds of images for each class such as:

<i>Z-line</i>	<i>Esophagitis</i>	<i>Pylorus</i>	<i>Polyps</i>
<p>The Z-line marks the transition site between the esophagus and the stomach.</p> 	<p>Esophagitis is an inflammation of the esophagus visible as a break in the esophageal mucosa in relation to the Z-line.</p> 	<p>The pylorus is defined as the area around the opening from the stomach into the first part of the small bowel (duodenum).</p> 	<p>Polyps are lesions within the bowel detectable as mucosal outgrowths.</p> 
<i>Cecum</i>	<i>Ulcerative Colitis</i>		
<p>The cecum is the most proximal part of the large bowel.</p> 	<p>Ulcerative colitis is a chronic inflammatory disease affecting the large bowel.</p> 		

First phase:

we used is a machine learning approach called support vector machine (SVM) that is Supervised learning is typically done in the context of classification, when we want to map input to output labels, or regression, when we want to map input to a continuous output, so we prepared the dataset before classify

Firstly by reading images with its different anatomical landmarks each one have positive and negative class, and it's for linearly separable binary set ,main Goal to design a hyper plane that classify all training vectors into two classes ,The best model that leaves the maximum margin from both classes. the two classes labels $+1$ (positive examples and -1 (negative examples).

Support Vector Machines

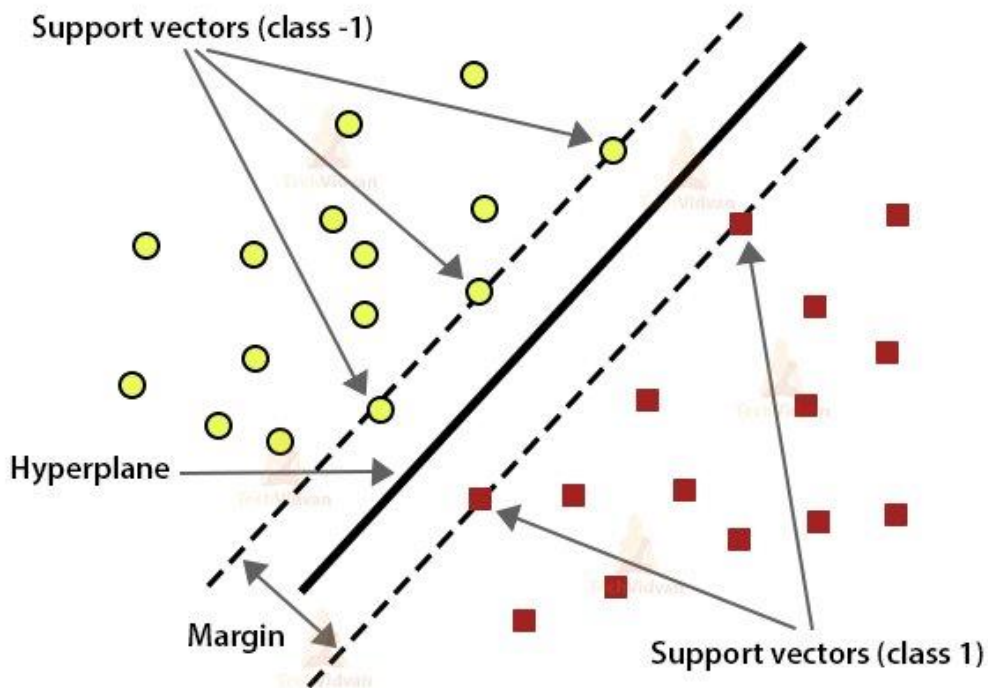


Fig 1.6

Secondly Preprocessing images

Removing the green square

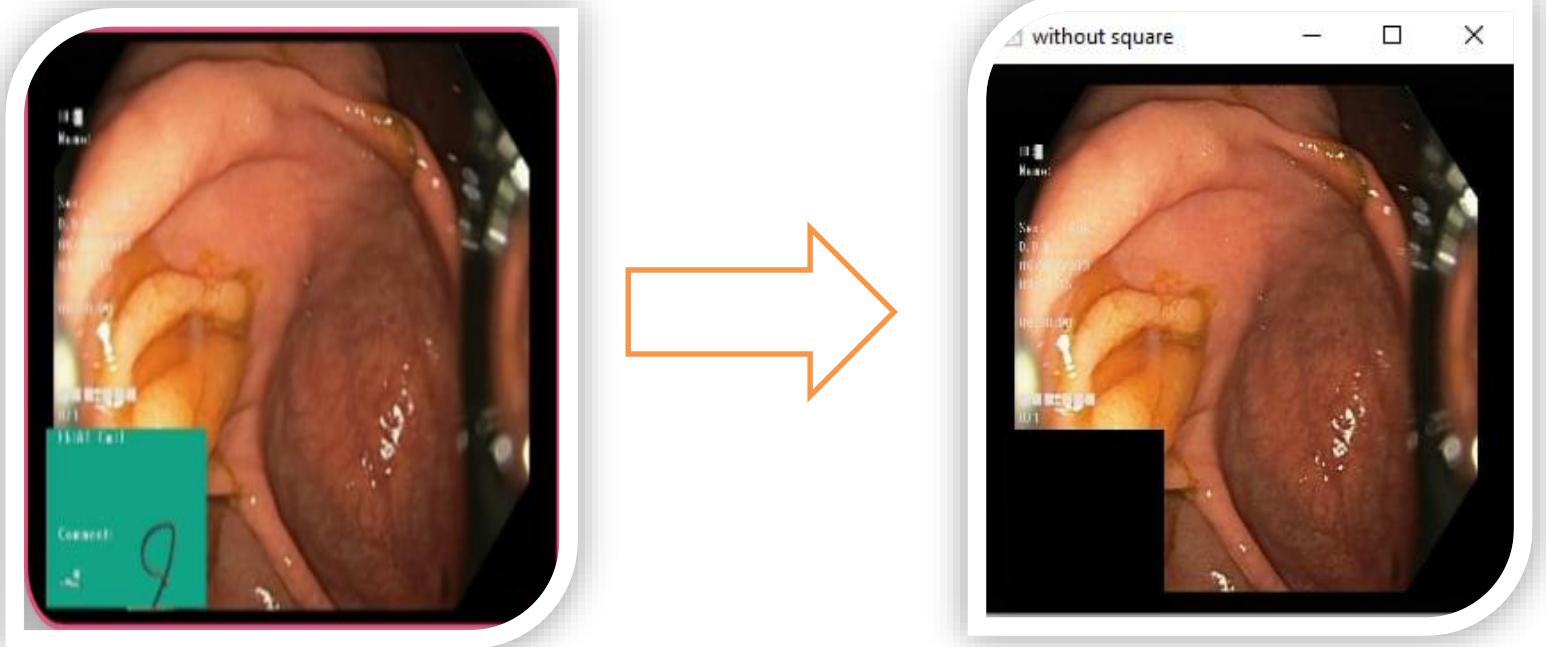
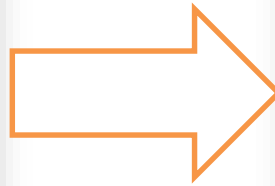


Fig 1.7

Blurring images with Gaussian Blur



Edge detection

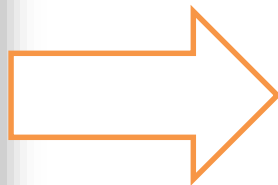


Fig 1.8

Finding contours



fig 1.9

Flatten images

Convert images into 1-d array

then the images begin to classify using SVM model and result in accuracy 87%

Second stage :

we used deep learning model called Convolutional Neural Networks(CNN) Deep learning is a subset of machine learning, which is essentially a neural network with three or more layers. These neural networks attempt to simulate the behavior of the human brain CNN, is a deep learning neural network designed for processing structured arrays of data such as images. Convolutional neural networks are widely used in computer vision and have become the state of the art for many visual applications such as image classification, and have also found success in natural language processing for text classification, so we perform preprocessing on data like we mentioned above and used different structures of CNN models. CNN is a feed-forward network that can extract topological properties from an image.

Like almost every other neural networks they are trained with a version of the back-propagation algorithm. CNN are designed to recognize

visual patterns directly from pixel images with minimal preprocessing. They can recognize patterns with extreme variability (such as handwritten characters).

CNN is a deep learning neural network designed for processing structured arrays of data such as images. CNN are widely used in computer vision and have become the state of the art for many visual applications such as image classification, and have also found success in natural language processing for text classification.

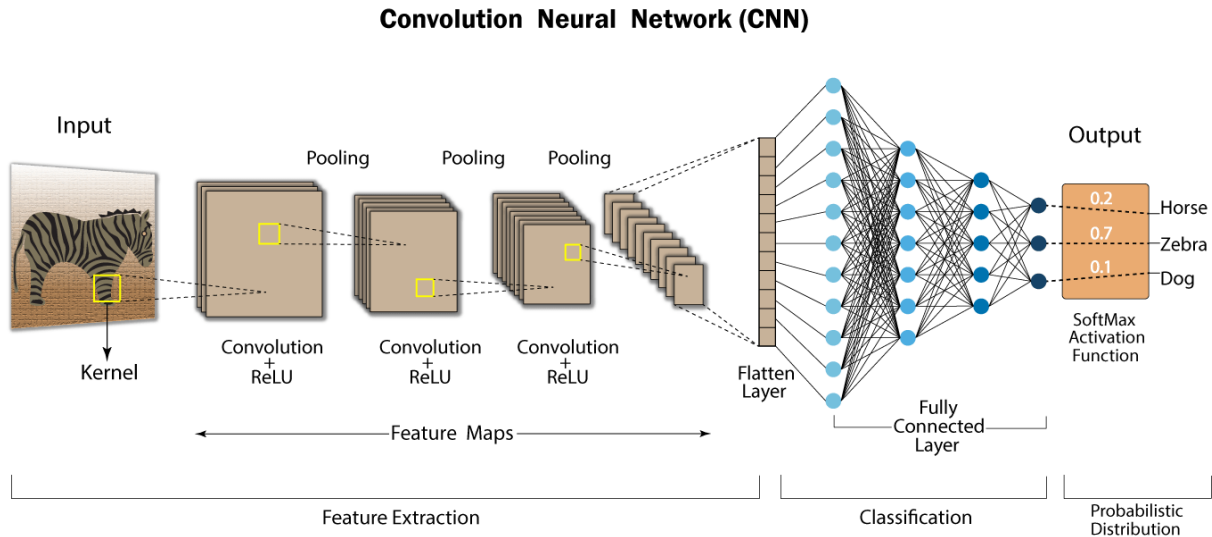


Fig 1.10

First we build a structure of CNN model from scratch, that contains conv2D with filter size 32 , max Pool 2*2, conv2D with filter size 64 and max Pool 2*2 , conv2D with filter size 128 and max Pool 2*2 ,

conv2D with filter size 256 and max Pool 2*2 , conv2D with filter size 512 and max Pool 2*2 then flatten and Dense, and we trained this model with the data that contains 6,000 images splitting it into 60% train data ,40% test data with 30 epochs.This model result in accuracy 73%

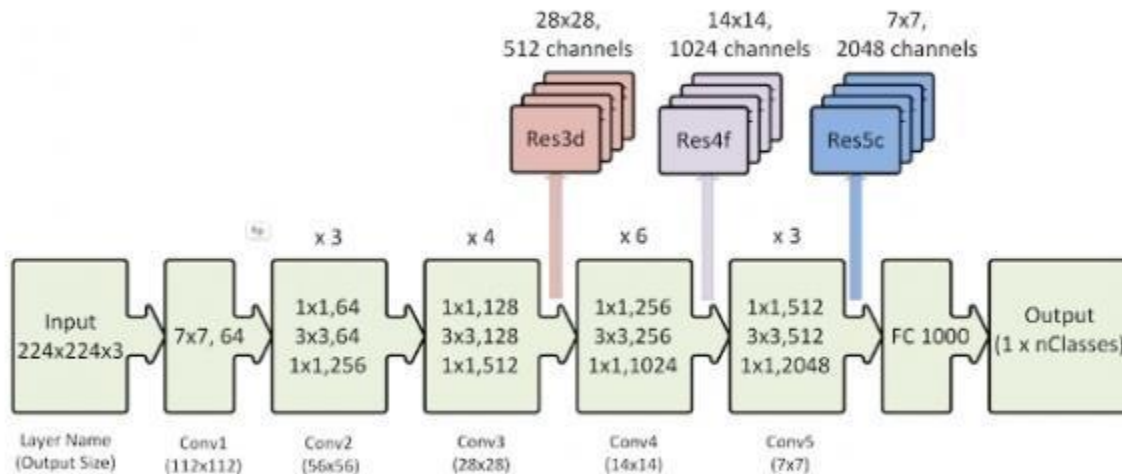


Fig1.11

For improving this accuracy we tried building another complex model with more layers and more number of filters: we build conv2D with filter size 200 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 240 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 280 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 280 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 320 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 360 ,maxPool 2*2 and Dropout with rate equals 0.2, conv2D with filter size 400 ,maxPool 2*2 and Dropout with rate equals 0.2 ,Flatten ,three layers of Dense with size 512,Dropout with rate equals 0.2 and Dense, to fit with data splitting the data also into 60% train ,20%validate and 20%test but with 70 epochs

and this model give accuracy 88%

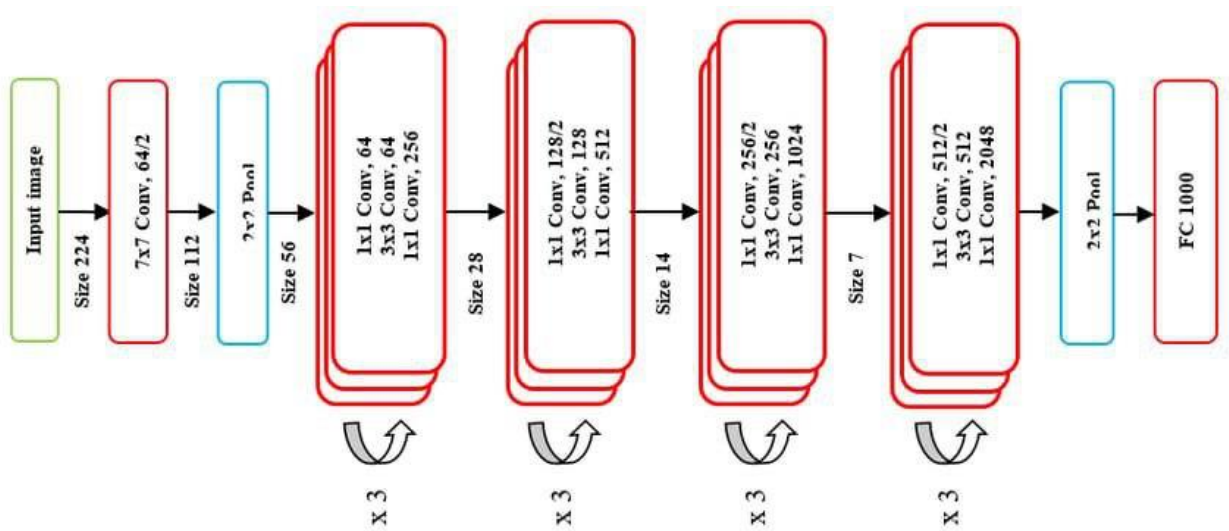


Fig1.12

We tried using built-in models in the hope of getting higher accuracy. First we used the built-in model VGG19 with 25 epochs and this model led to accuracy 87%.

Finally:

we used built-in ResNet50 hoping of getting a high accuracy with 25 epochs and this model led to accuracy 93%.

1.6 The used tools in the project

The Software tools that used in project:

Operating system	Applications
<ul style="list-style-type: none">• Mac• Windows	<ul style="list-style-type: none">• Visual studio Code• HTML• PHP• CSS• JS• PyCharm• Python 3.10• UsbServer• MAMPsServer• PHPMyAdmin• Skitlearn• Opencv• Tensorflow• Keras

1.7 Report Organization

Chapter Two: Related Work

In chapter two, we will establish other work associated with our research. There, we show that different ways to achieve our goals. We will declare the authors of these methods, the classification method they use in their project, and off course the accuracy.

Chapter Three: System Analysis

In chapter three, we will talk about project specification. It will contain descriptions of how the program will be used from a user perspective and performance details such as usability, reliability, and stability. In addition, illustrate a use case diagram that emphasizes our program.

Chapter Four: System Design

The System Design Document describes the system requirements, operating environment, system and subsystem architecture, files and database design, input formats, output layouts, human-machine interfaces, detailed design, processing logic, and external interfaces using some diagrams. System Component Diagram, System Class Diagrams, Sequence Diagrams, Project ERD, System GUI Design

Chapter Five: Implementation and Testing

This phase includes screen shots of training and testing of (SVM,) model, CNN model

Screenshots of the system running and samples of the applied test cases.

Chapter Two

Related work

Colorectal Cancer Detection Based on Deep Learning :

The traditional idea:

was for a specialized pathologist to make an assessment of (H&E)-stained tissue sections by microscopy of multiple samples taken from one patient

This is step need more attention to detail .

In addition diagnoses depended on a individual pathologist's experience

Solution: using CNN model and using digitized H&E-stained histology slides

Accuracy is 89%

Dataset:

307 (85 normal & 222 colorectal cancer tissue)

275 slides (76 normal and 199 containing colorectal cancer) were randomly selected for training; the remaining slides were used for testing (9 normal and 23 colorectal cancer)

Work limitations:

-the preparation needed to build such model, from samples taken from patients' colon (health risk) to digitizing the slides is intensive and time consuming -Data amount is very small

Deep Learning in Image Classification using ResNet **Variants for Detection of Colorectal Cancer**

The traditional idea:

A pathologist collects biopsies

From all parts of colon, cells to identify cancer

They examine the tissue so that they can tell what stage this cancer is in.

This is done by examining the morphology of the cells and observing the changes that occur in them, as well as in the components of tissues visually under a microscope.

However, pathologists often disagree with regard to classifications of cancer, it is not enough to rely on the expertise of scientists alone.

Solution: Medical imaging presently serves to distinguish the diseases in various components of the body. And 3 models of deep learning (ResNet18, ResNet50, ResNet18&ResNet50)

Accuracy is 85%

Dataset:

-The data consist of 165 images with 74 benign tumors images and 91 malignant tumors images.

-This data was attained with Scanner

Work limitations

ResNet 50 take more time than ResNet 18 to deal with large data
Data amount is very small to depend on to prepare model

Work Differences :

- our project used CNN model and SVM model, but project 1 used only CNN model , project 2 used 3 models
ResNet18,ResNet50,(ResNet18&ResNet50)
- in our project have 8000 images on another hand project 1 have 307 images, project 2 have 165 images [very small data]
- our data set used colonoscopy video , other projects used slides digitized from clinical samples by Scanner

Chapter Three

System Analysis

3.1 Project specification

3.1.1. Functional requirement

- Registration: If this is the first time for the user to use the website, he/she will register.
- Log in: If the user already registers before, the user will log in to the app with his user's name and password
- profile: the user can show his history, and he could back to the history and check the results and accuracy.
- Adding information for new Patient Or choose existing one
- Upload image as input
- image processing
- Determine if this image have cancer or not by model
- Output the classification and accuracy.

3.1.2. Non-functional requirement

- Usability
The site is easy for use
- Security
Login Authentication
The site is secure, because no one can access without e-mail

- **Reusability**
The system can be used in another medical purpose without a lot of change.
- **Response Time**
The site doesn't take more than 5 second to classify the images
- **Robustness**
The system can deal with errors from the user during execution without crashing.
The deployed model doesn't suffer from overfitting or underfitting. The difference between train and test accuracy less than 1%.
- **Availability**
The site is available 24 hours
- **Maintainability**
It's easy to deploy new model better than the model used in app.
The system can be easily repaired and maintained

3.2. Use case Diagrams

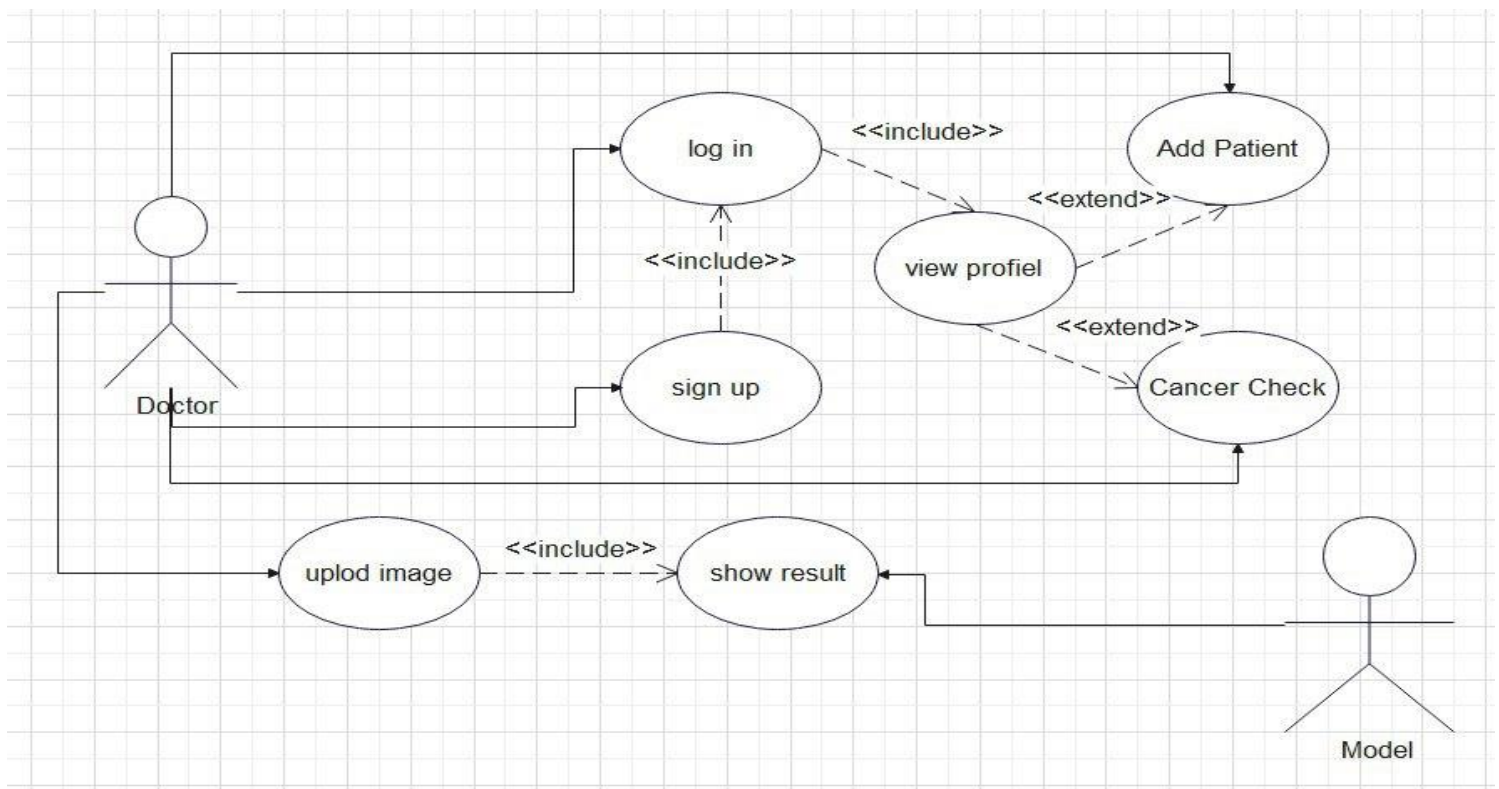


Fig 3.1

Chapter Four

System Design

System Component Diagram

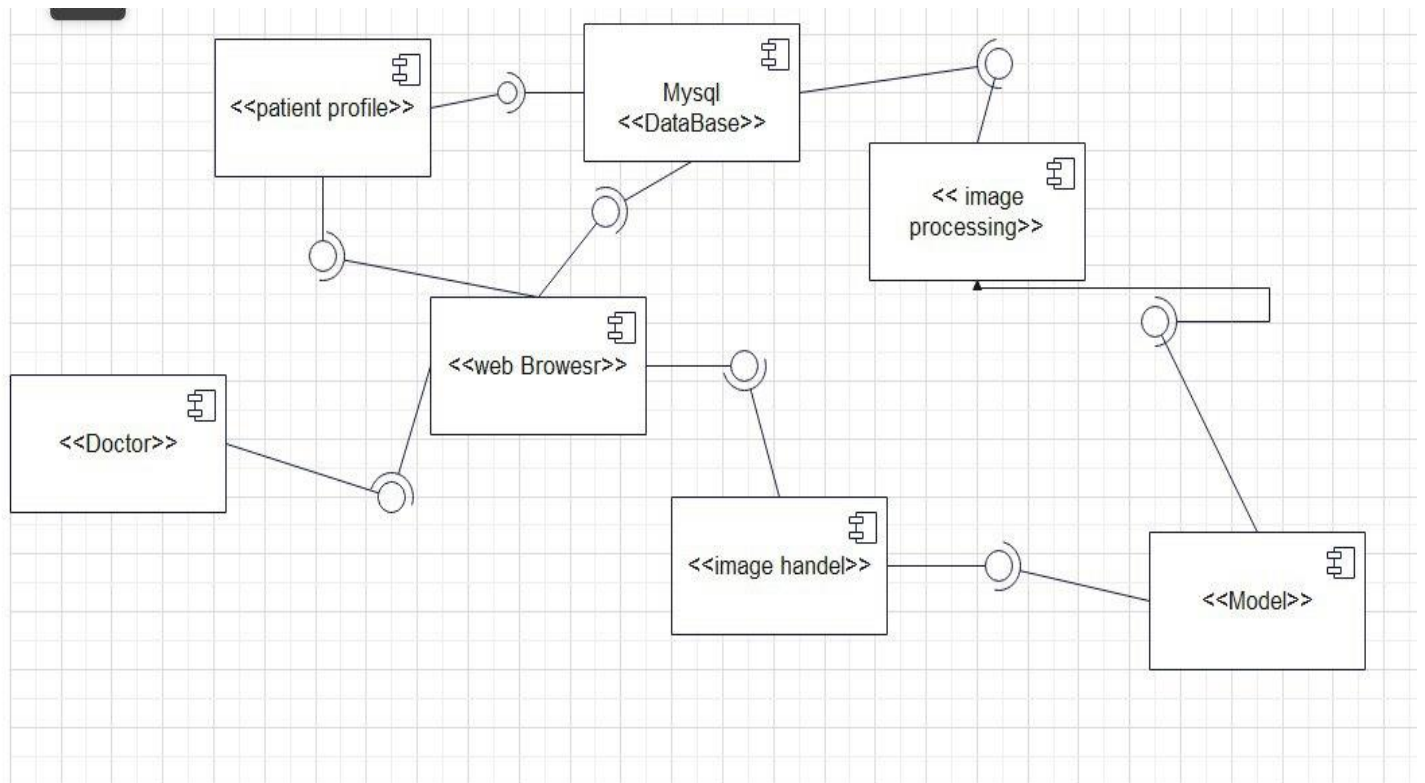


Fig 4.1

System class Diagram

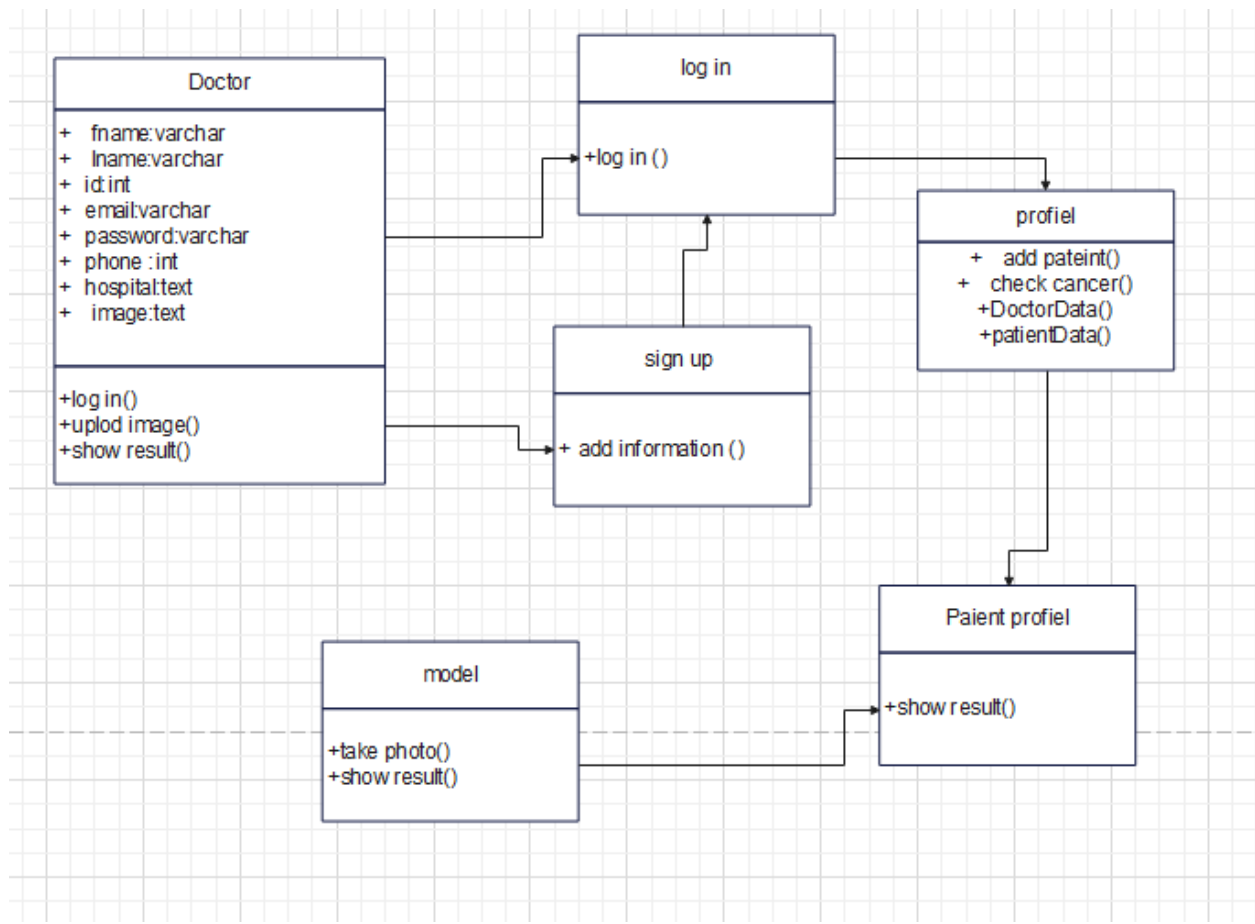


Fig 4.2

Sequence Diagrams

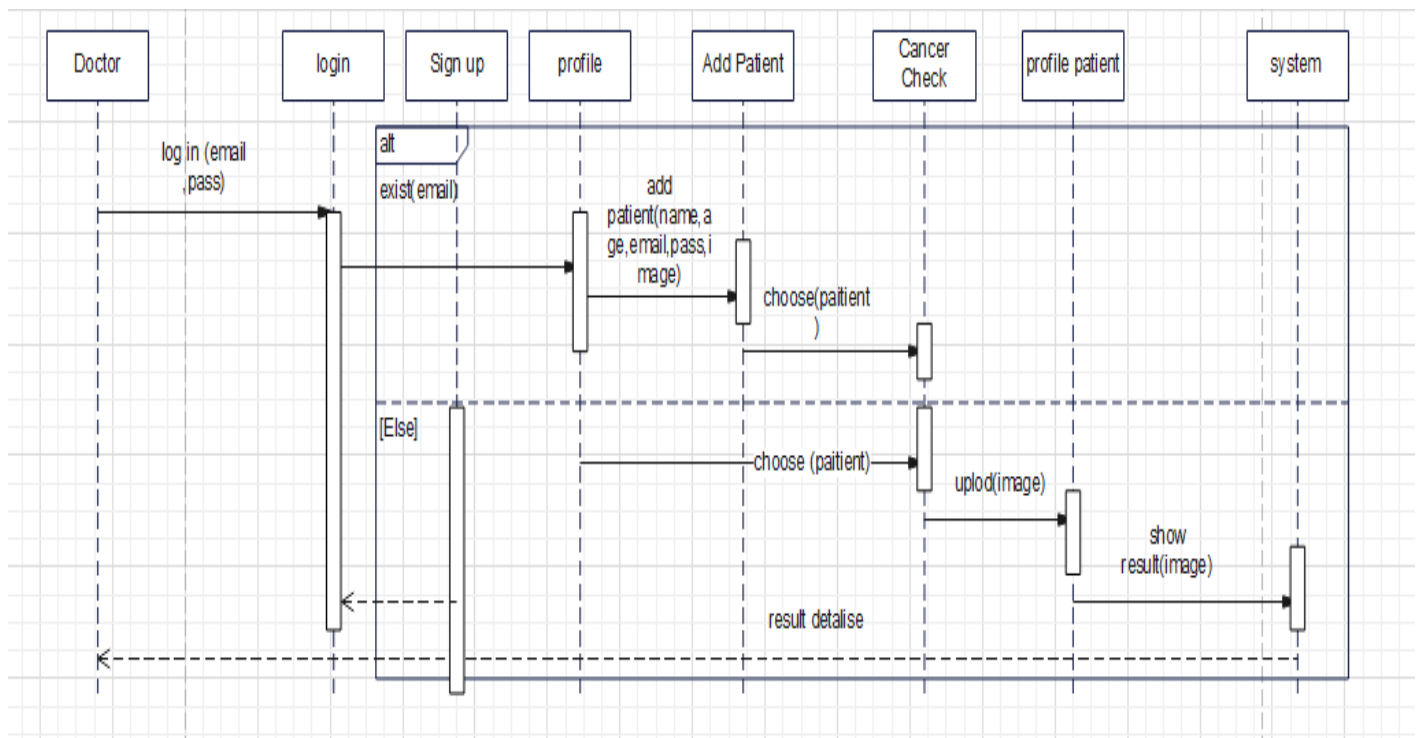


Fig 4.3

Chapter Five

Implementation and Testing

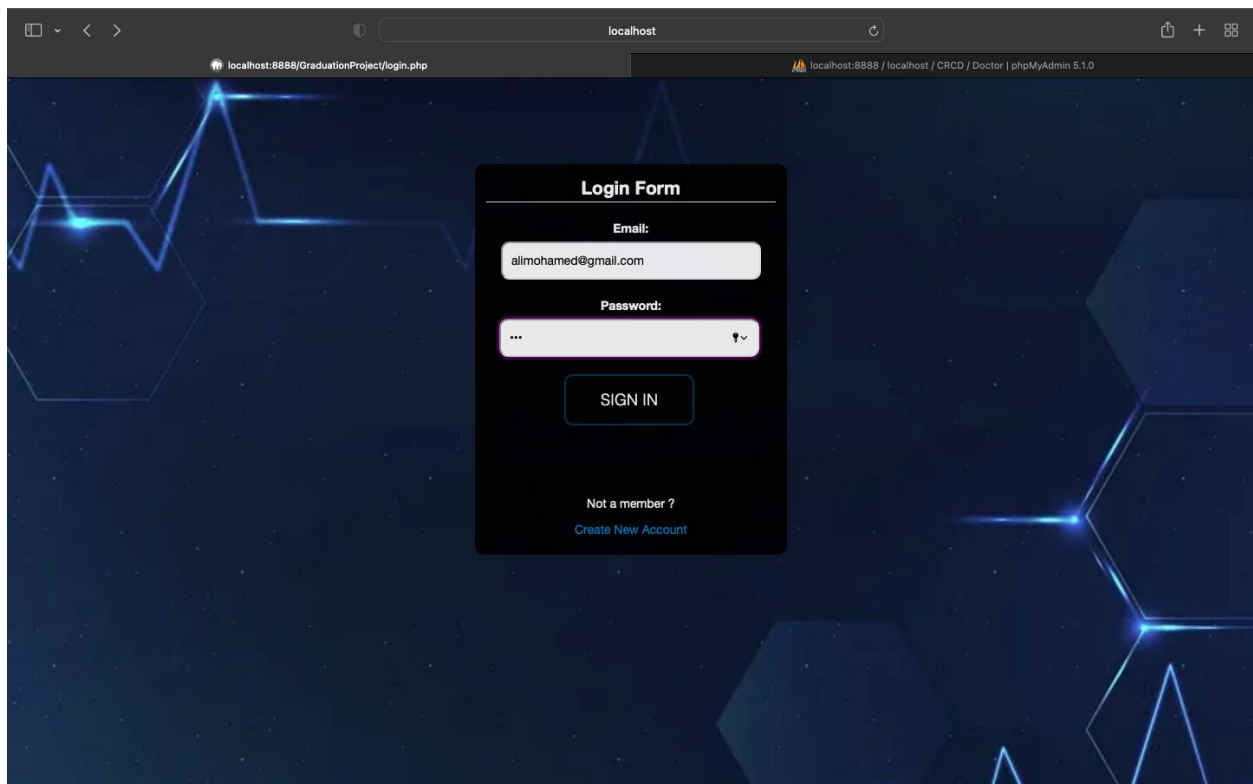
Register Form

The screenshot displays a web browser window with the address bar showing 'localhost'. The page title is 'Register Form'. The form is set against a dark blue background with glowing blue hexagonal patterns. The form fields are as follows:

- First Name ***: Input field containing 'Ali'.
- Last Name ***: Input field containing 'Mohamed'.
- Email ***: Input field containing 'alimohamed@gmail.com'.
- Password ***: Input field with masked characters '....' and a toggle icon.
- phone**: Input field containing '01144117828'.
- Hospital**: Input field containing '57357'.
- Add Image**: A section with a 'Choose File' button and the filename 'Ali.jpg'.

At the bottom of the form are two buttons: 'RESET' and 'SIGN UP'.

Log in Page



localhost

localhost:8888/GraduationProject/login.php

localhost:8888 / localhost / CRCD / Doctor | phpMyAdmin 5.1.0

Login Form

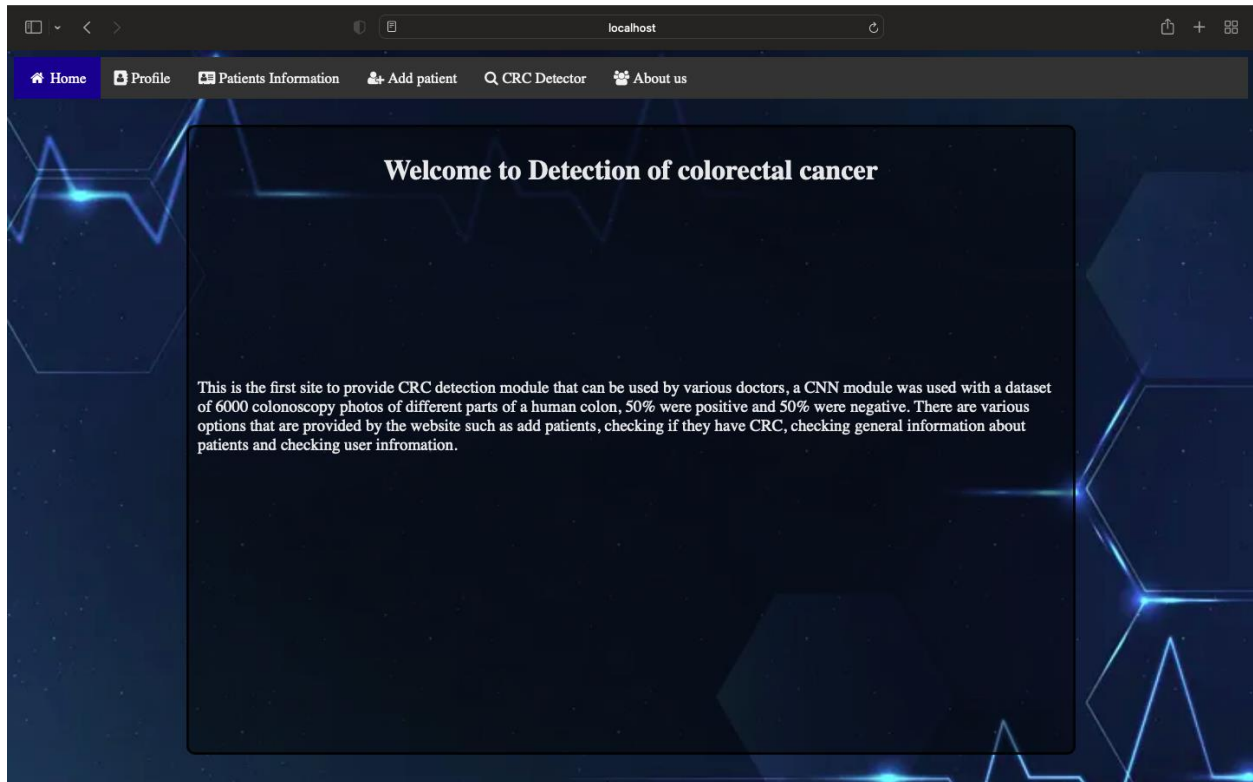
Email:
alimohamed@gmail.com

Password:
...

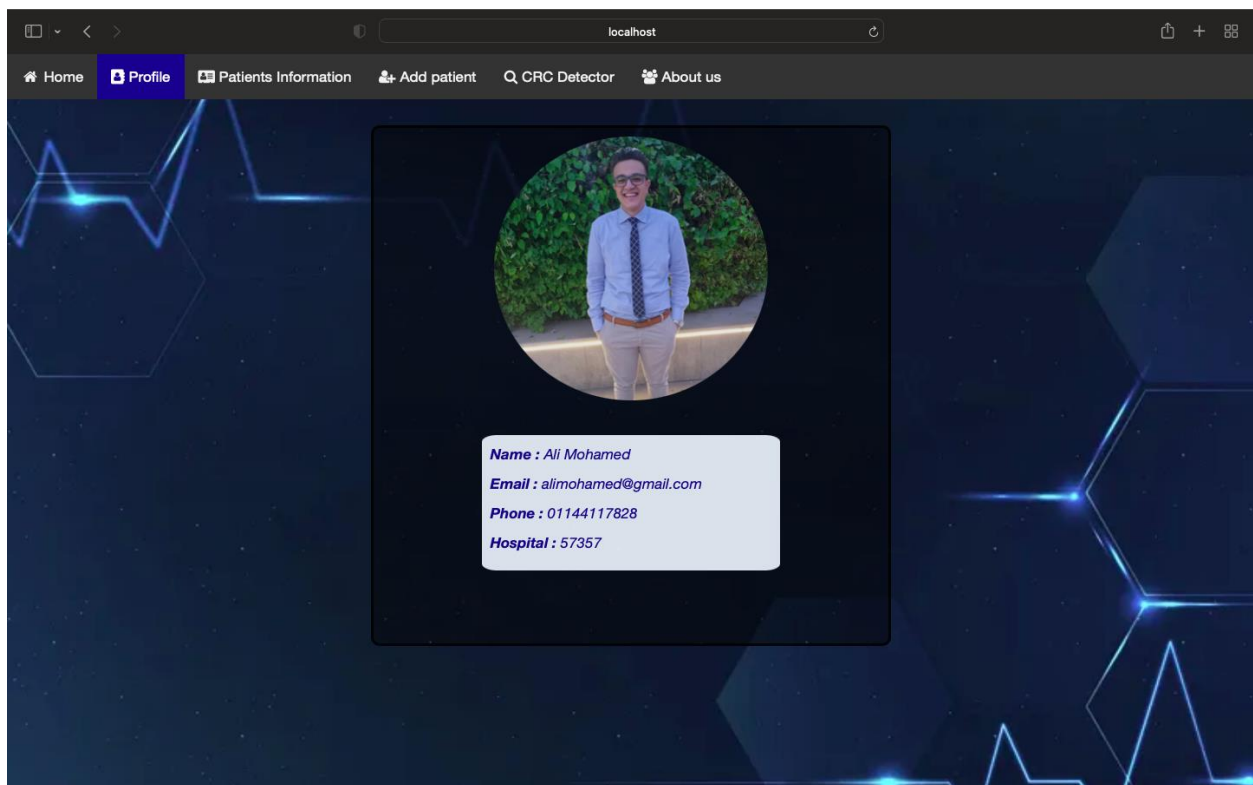
SIGN IN

Not a member ?
[Create New Account](#)

Home



Profile



Add Patient

The screenshot shows a web browser window with the URL 'localhost'. The navigation bar includes links for Home, Profile, Patients Information, Add patient (highlighted), CRC Detector, and About us. The main content area features a 'Patient Form' with the following fields:

- Patient Name ***: A text input field containing 'Said Mohamed'.
- Age ***: A numeric input field containing '25'.
- Medical History**: A text area containing 'No past diseases'.
- Patient Image (If available)**: A section with a 'Choose File' button and a file name '0d7bb96c-ea11...ab2f76ca.jpg'.

At the bottom of the form are two buttons: 'RESET' and 'ADD PATIENT'.

This screenshot shows the same 'Add Patient' form, but with a 'Choose Files to Upload' dialog box open over the 'Patient Image' section. The dialog box displays a list of files in the 'Downloads' folder, with 'patient.jpg' selected. A preview of the selected image is shown on the right side of the dialog. The file information indicates it is a 'JPEG image - 2 KB'.

The background form is partially visible, showing the 'Patient Form' title and the 'Patient Image (If available)' section with a 'Choose File' button and the text 'no file selected'.

Data Base inserted in Patient table

The screenshot displays the phpMyAdmin web interface. The left sidebar shows the database structure with 'CRCD' selected. The main panel shows the 'Patient' table with a single row of data. The table has columns: id, name, age, medical_history, patient_image, location_in_colon, image, CRC, and doctor_id. The row contains the following data: id 1, name Said Mohamed, age 25, medical_history No past diseases, patient_image 0d7bb96c-ea11-4dbe-98c0-4ddaab2f76ca.jpg, location_in_colon NULL, image NULL, CRC NULL, and doctor_id 5. The interface includes a top navigation bar, a left sidebar, and a main content area with a table view and a console at the bottom.

Showing rows 0 - 0 (1 total, Query took 0.0004 seconds.)

```
SELECT * FROM `Patient`
```

Profiling [Edit inline] [Edit] [Explain SQL] [Create PHP code] [Refresh]

Show all | Number of rows: 25 | Filter rows: Search this table

+ Options

	id	name	age	medical_history	patient_image	location_in_colon	image	CRC	doctor_id
<input type="checkbox"/>	1	Said Mohamed	25	No past diseases	0d7bb96c-ea11-4dbe-98c0-4ddaab2f76ca.jpg	NULL	NULL	NULL	5

Check all | With selected: Edit Copy Delete Export

Show all | Number of rows: 25 | Filter rows: Search this table

Query results operations

Print Copy to clipboard Export Display chart Create view

CRC Detector

1.

The screenshot shows a web browser at localhost with the CRC Detector application. The navigation bar includes Home, Profile, Patients Information, Add patient, CRC Detector (active), and About us. The main content area features a 'Check for cancer' form with the following elements:

- Select patient:** A dropdown menu showing 'Said Mohamed'.
- Select Location in colon:** Three radio buttons on the left and three labels on the right: 'Z_line', 'Polyps', and 'Cecum'.
- Patient Colon Image:** A 'Choose File' button and the text 'no file selected'.
- Buttons:** 'RESET' and 'RESULT' buttons at the bottom.

2.

This screenshot shows the same CRC Detector web application, but with the 'Select patient' dropdown menu open. The dropdown list displays two options: 'Said Mohamed' (which is highlighted with a checkmark) and 'Essam Ali'.

The rest of the form, including the 'Select Location in colon' section with radio buttons and labels ('Z_line', 'Polyps', 'Cecum'), the 'Patient Colon Image' section with the 'Choose File' button and 'no file selected' text, and the 'RESET' and 'RESULT' buttons, remains the same as in the first screenshot.

3.

The screenshot displays a web application interface for a CRC Detector. The browser address bar shows 'localhost'. The application has a dark theme with a navigation bar at the top containing links: Home, Profile, Patients Information, Add patient, CRC Detector (active), and About us. The main content area features a 'Check for cancer' modal form. This form includes a 'Select patient' dropdown menu with 'Said Mohamed' selected, a 'Select Location in colon' section with three radio buttons (the first is selected), and a 'Patient Colon Image' section with a 'Choose File' button and the filename 'cecum.jpg'. At the bottom of the modal are 'RESET' and 'RESULT' buttons. The background of the application has a dark blue space-themed pattern with glowing geometric shapes.


Patient Result

localhost

localhost:8888 / localhost / CRCDD / Patient | phpMyAdmin 5.1.0

Home Profile Patients Information Add patient CRC Detector About us

Patient Results



Patient Name


cecum

Medical History

No past diseases

Colon Image

666



localhost

localhost:8888 / localhost / CRCDD / Patient | phpMyAdmin 5.1.0

cecum

Medical History

No past diseases

Colon Image

666

11/06/2015
09:34:57
0.35-1
0.21

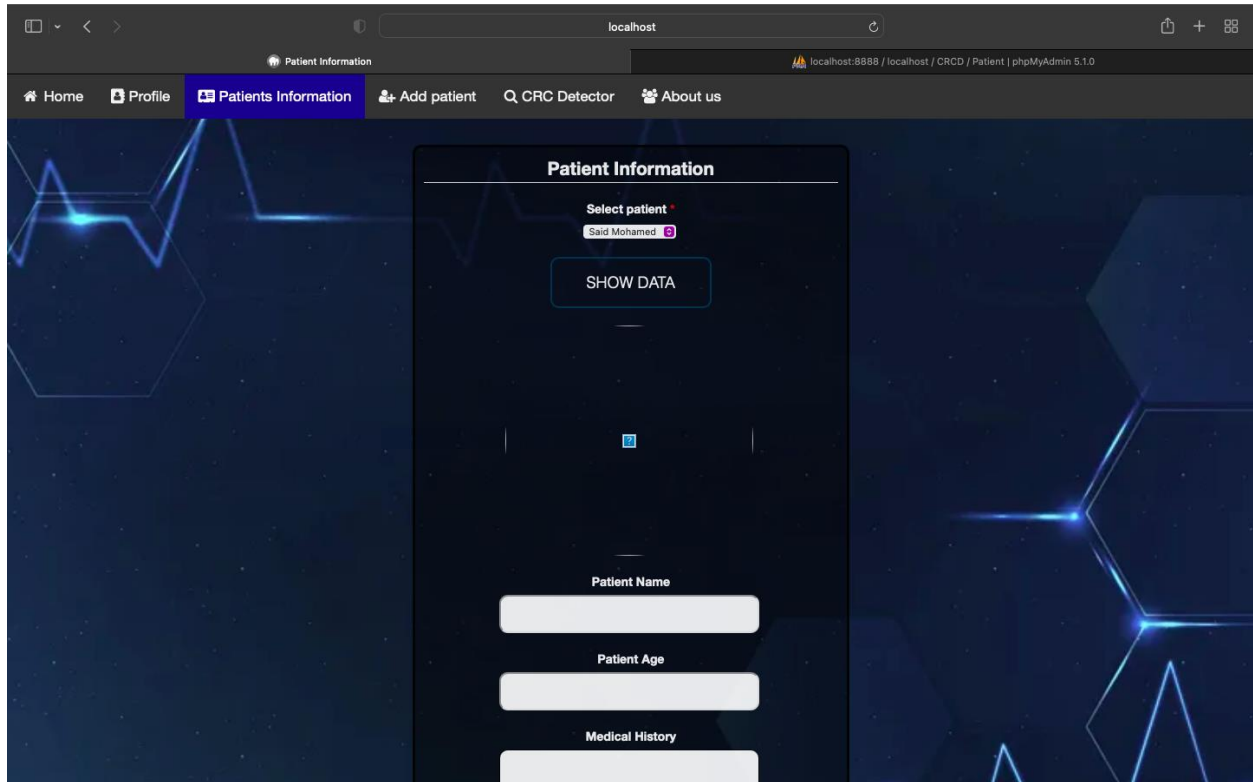
Image Location

cecum

CRC

Negative

Patient Information



localhost

Patient Information

localhost:8888 / localhost / CRCD / Patient | phpMyAdmin 5.1.0

Home Profile Patients Information Add patient CRC Detector About us

Patient Information

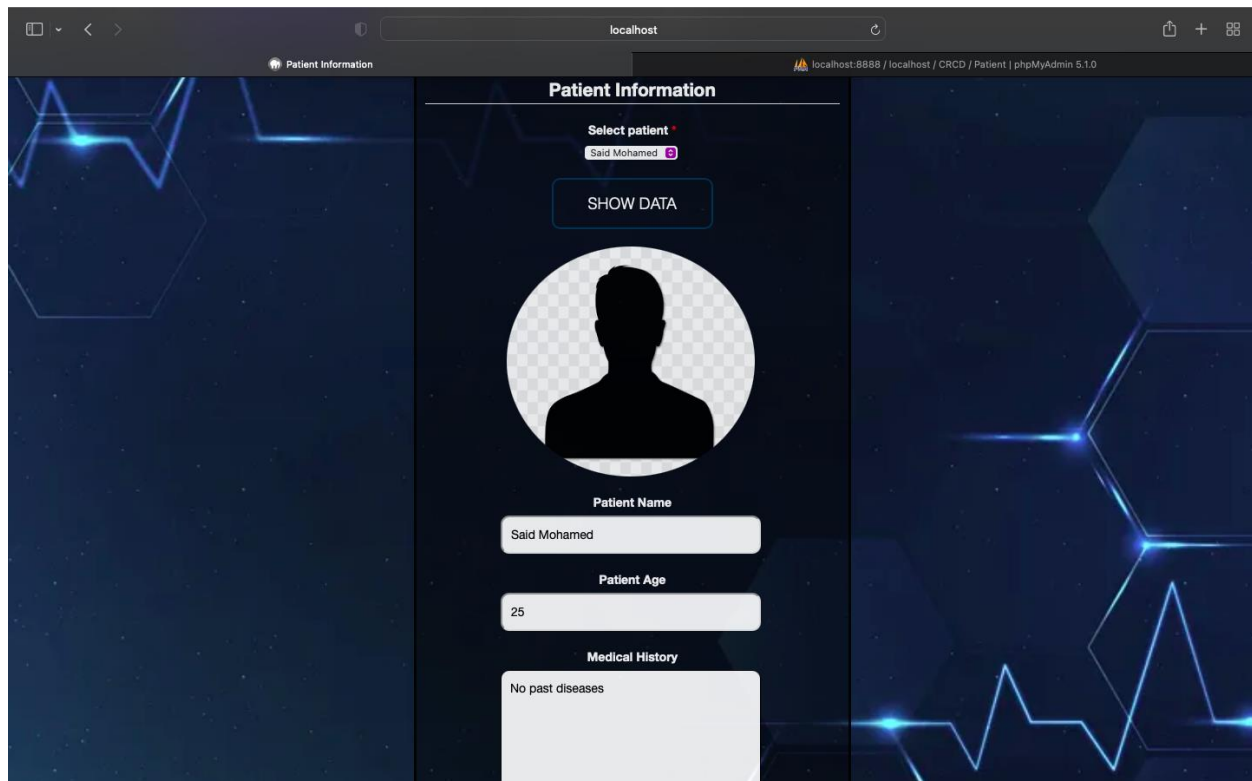
Select patient
Said Mohamed

SHOW DATA

Patient Name

Patient Age

Medical History



localhost

Patient Information

localhost:8888 / localhost / CRCD / Patient | phpMyAdmin 5.1.0

Home Profile Patients Information Add patient CRC Detector About us

Patient Information

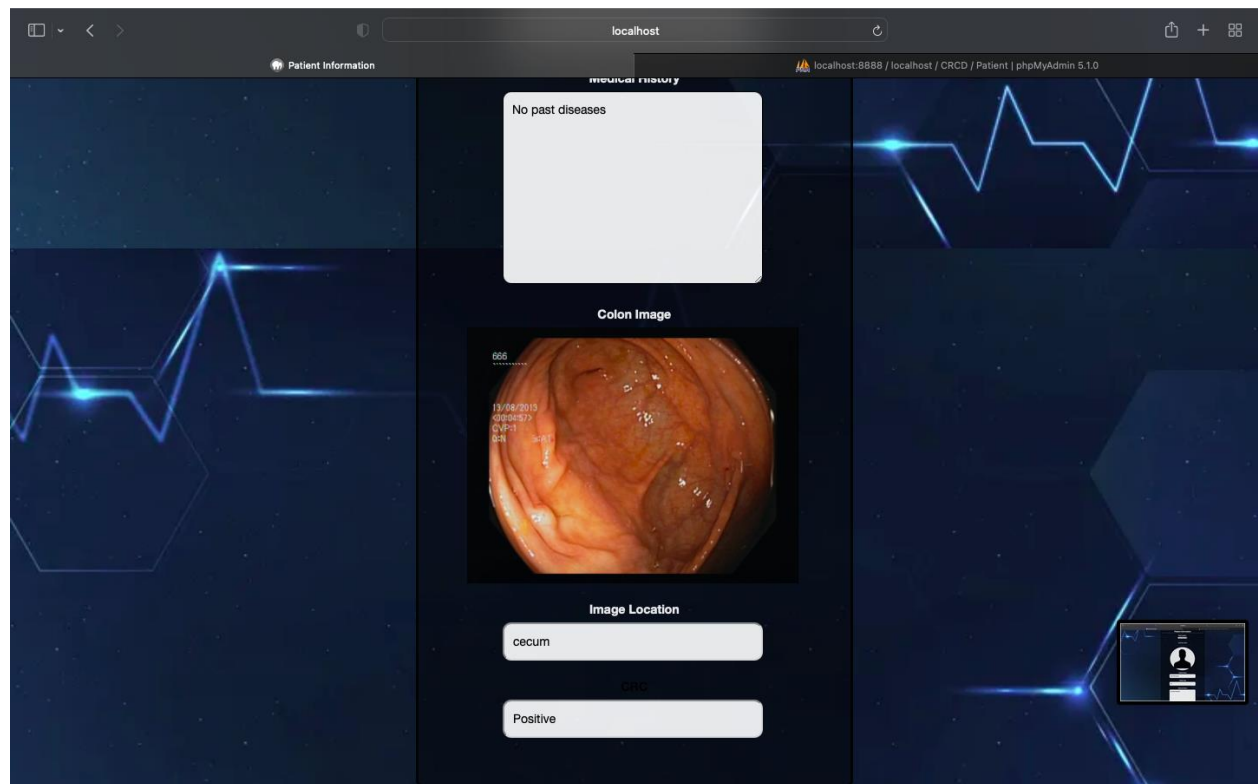
Select patient
Said Mohamed

SHOW DATA

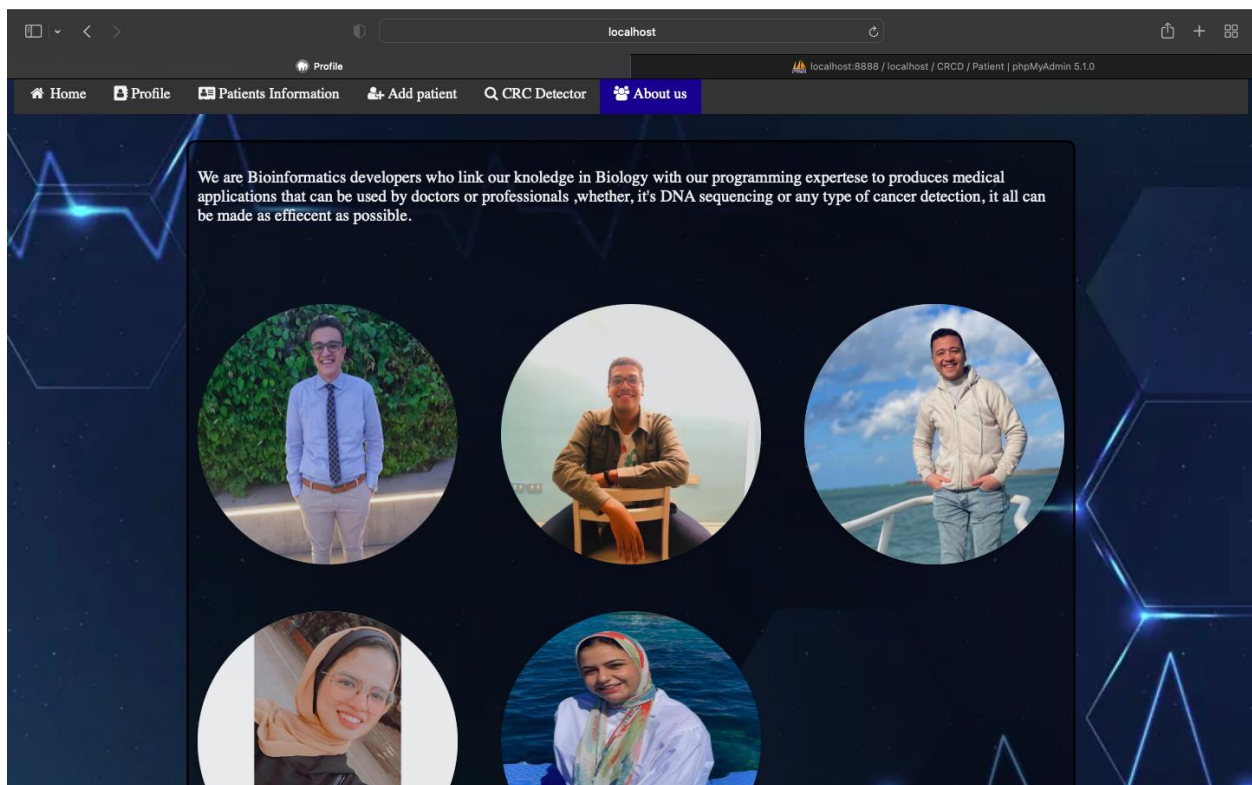
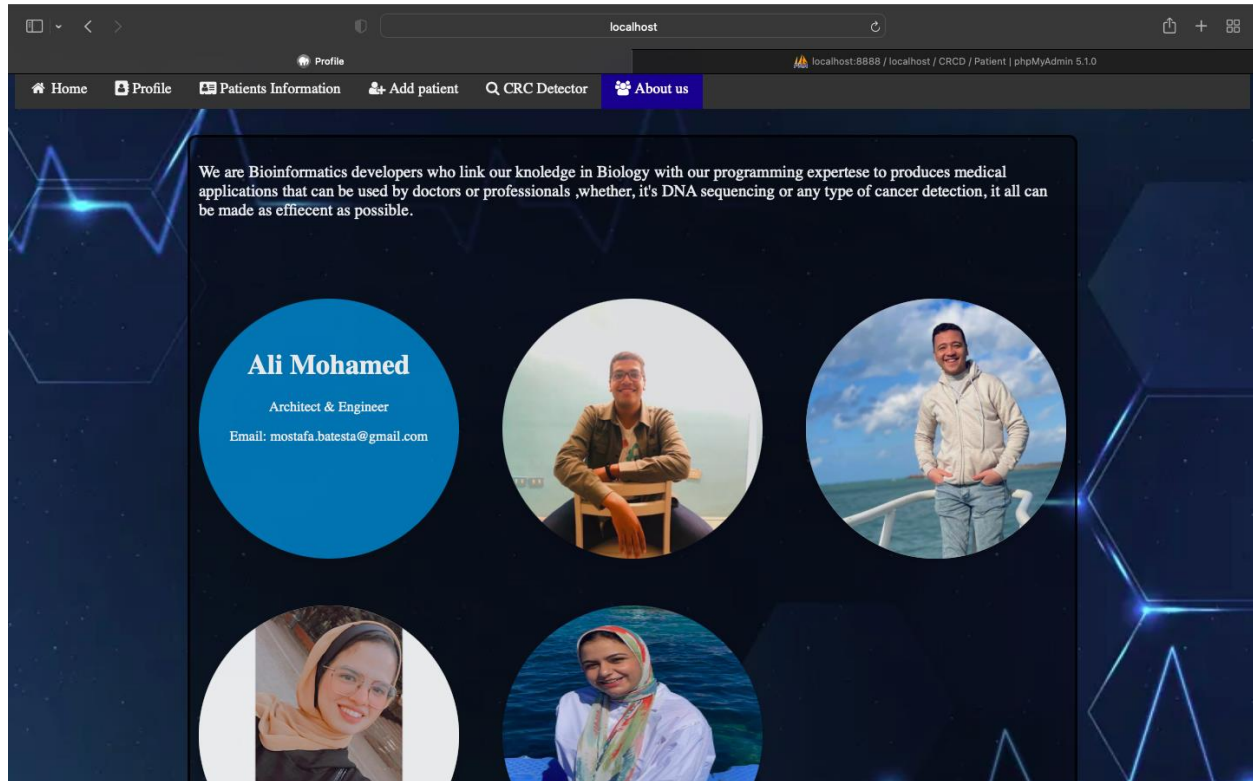
Patient Name
Said Mohamed

Patient Age
25

Medical History
No past diseases



About Us



Database Schema

localhost:8888/GraduationProject/login.php

localhost:8888 / localhost / CRCD | phpMyAdmin 5.1.0

Launch Meeting - Zoom

phpMyAdmin

Recent Favorites

New

CRCD

New

Doctor

Patient

information_schema

mysql

performance_schema

sys

Filters

Containing the word:

Table	Action	Rows	Type	Collation	Size	Overhead
<input type="checkbox"/> Doctor	Browse Structure Search Insert Empty Drop	2	InnoDB	utf8_general_ci	32.0 KiB	-
<input type="checkbox"/> Patient	Browse Structure Search Insert Empty Drop	2	InnoDB	utf8_general_ci	32.0 KiB	-
2 tables	Sum	4	InnoDB	utf8_general_ci	64.0 KiB	0 B

☐ Check all With selected:

[Print](#) [Data dictionary](#)

[Create table](#)

Name: Number of columns: 4

Go

Console

Tables

The screenshot shows the phpMyAdmin interface for the 'Doctor' table in the 'CRCD' database. The table structure is as follows:

#	Name	Type	Collation	Attributes	Null	Default	Comments	Extra	Action
1	id	int(11)			No	None		AUTO_INCREMENT	Change Drop More
2	fname	varchar(30)	utf8_general_ci		No	None			Change Drop More
3	lname	varchar(100)	utf8_general_ci		No	None			Change Drop More
4	email	varchar(200)	utf8_general_ci		No	None			Change Drop More
5	password	varchar(100)	utf8_general_ci		No	None			Change Drop More
6	phone	varchar(11)	utf8_general_ci		No	None			Change Drop More
7	hospital	varchar(100)	utf8_general_ci		No	None			Change Drop More
8	image	varchar(200)	utf8_general_ci		Yes	NULL			Change Drop More

Below the table structure, the 'Indexes' section shows two indexes:

Action	Keyname	Type	Unique	Packed	Column	Cardinality	Collation	Null	Comment
Edit Rename Drop	PRIMARY	BTREE	Yes	No	id	2	A	No	
Edit Rename Drop	email	BTREE	Yes	No	email	2	A	No	

The 'Partitions' section shows 'No partitioning defined!'.

The screenshot shows the phpMyAdmin interface for the 'Patient' table in the 'CRCD' database. The table structure is as follows:

#	Name	Type	Collation	Attributes	Null	Default	Comments	Extra	Action
1	id	int(11)			No	None		AUTO_INCREMENT	Change Drop More
2	name	varchar(100)	utf8_general_ci		No	None			Change Drop More
3	age	int(11)			No	None			Change Drop More
4	medical_history	varchar(1000)	utf8_general_ci		No	None			Change Drop More
5	patient_image	varchar(200)	utf8_general_ci		Yes	NULL			Change Drop More
6	location_in_colon	varchar(100)	utf8_general_ci		Yes	NULL			Change Drop More
7	Image	varchar(200)	utf8_general_ci		Yes	NULL			Change Drop More
8	CRC	tinyint(1)			Yes	NULL			Change Drop More
9	doctor_id	int(11)			No	None			Change Drop More

Below the table structure, the 'Indexes' section shows two indexes:

Action	Keyname	Type	Unique	Packed	Column	Cardinality	Collation	Null	Comment
Edit Rename Drop	PRIMARY	BTREE	Yes	No	id	2	A	No	
Edit Rename Drop	doctor_id	BTREE	No	No	doctor_id	1	A	No	

The 'Partitions' section shows 'No partitioning defined!'.

Appendix

Phase one model using supervised technique (SVM) with 87% accuracy

```
def svm(svm_x_train, svm_y_train, svm_x_test, svm_y_test):  
    svc_model = SVC(C=.1, kernel='poly', gamma=1)  
    svc_model.fit(svm_x_train, svm_y_train)  
    prediction = svc_model.predict(svm_x_test)  
  
    filename = 'finalized_model.sav'  
    pickle.dump(svc_model, open(filename, 'wb'))  
    return accuracy_score(svm_y_test, prediction) * 100
```

Phase two models using unsupervised technique (CNN)

First model with 73% accuracy

```
KerasModel = keras.models.Sequential([  
    keras.layers.Conv2D(32, kernel_size=(3, 3), input_shape=(size, size, 1)),  
    keras.layers.MaxPool2D(2, 2),  
    keras.layers.Conv2D(64, kernel_size=(3, 3)),  
    keras.layers.MaxPool2D(2, 2),  
    keras.layers.Conv2D(128, kernel_size=(3, 3)),  
    keras.layers.MaxPool2D(2, 2),  
    keras.layers.Conv2D(256, kernel_size=(5, 5)),  
    keras.layers.MaxPool2D(2, 2),  
    keras.layers.Conv2D(512, kernel_size=(5, 5)),  
    keras.layers.MaxPool2D(2, 2),  
    keras.layers.Flatten(),  
    keras.layers.Dense(6, activation='softmax')  
)
```

Second model with 88% accuracy

```
CNN_model.py - keras_CNN_model.py
KerasModel = keras.models.Sequential([
    keras.layers.Conv2D(200, kernel_size=(3, 3), activation='relu', input_shape=(size, size, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),

    keras.layers.Conv2D(240, kernel_size=(3, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),

    keras.layers.Conv2D(280, kernel_size=(3, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),

    keras.layers.Conv2D(320, kernel_size=(3, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),

    keras.layers.Conv2D(360, kernel_size=(3, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),
```

```
    keras.layers.Conv2D(400, kernel_size=(3, 3)),
    keras.layers.MaxPool2D(2, 2),
    keras.layers.Dropout(rate=0.2),

    keras.layers.Flatten(),
    keras.layers.Dense(512, activation='relu'),
    keras.layers.Dense(512, activation='relu'),
    keras.layers.Dense(512, activation='relu'),

    keras.layers.Dropout(rate=0.2),

    keras.layers.Dense(6, activation='softmax', kernel_regularizer=l2(0.01),
                        bias_regularizer=l2(0.01))
])
```

Third model CNN VGG19 with 87% accuracy

```
inputs = Input(shape=(224,224,3))
base_model = VGG19(include_top=False, weights='imagenet', input_shape=(224, 224,3))
base_model.trainable = False

added_layers = base_model(inputs, training=False)
added_layers = Flatten()(added_layers)
added_layers = Dense(100,activation='relu')(added_layers)
added_layers = Dense(50, activation='relu')(added_layers)
final_layer = Dense(6, activation='softmax')(added_layers)

KerasModel = Model(inputs, final_layer)
KerasModel.compile(optimizer='adam', loss='categorical_crossentropy', metrics=['accuracy'])
epochs = 25
KerasModel.fit(x_train, to_categorical(y_train), epochs=epochs, batch_size=64, verbose=1,
                validation_data=(x_valid, to_categorical(y_valid)))
```

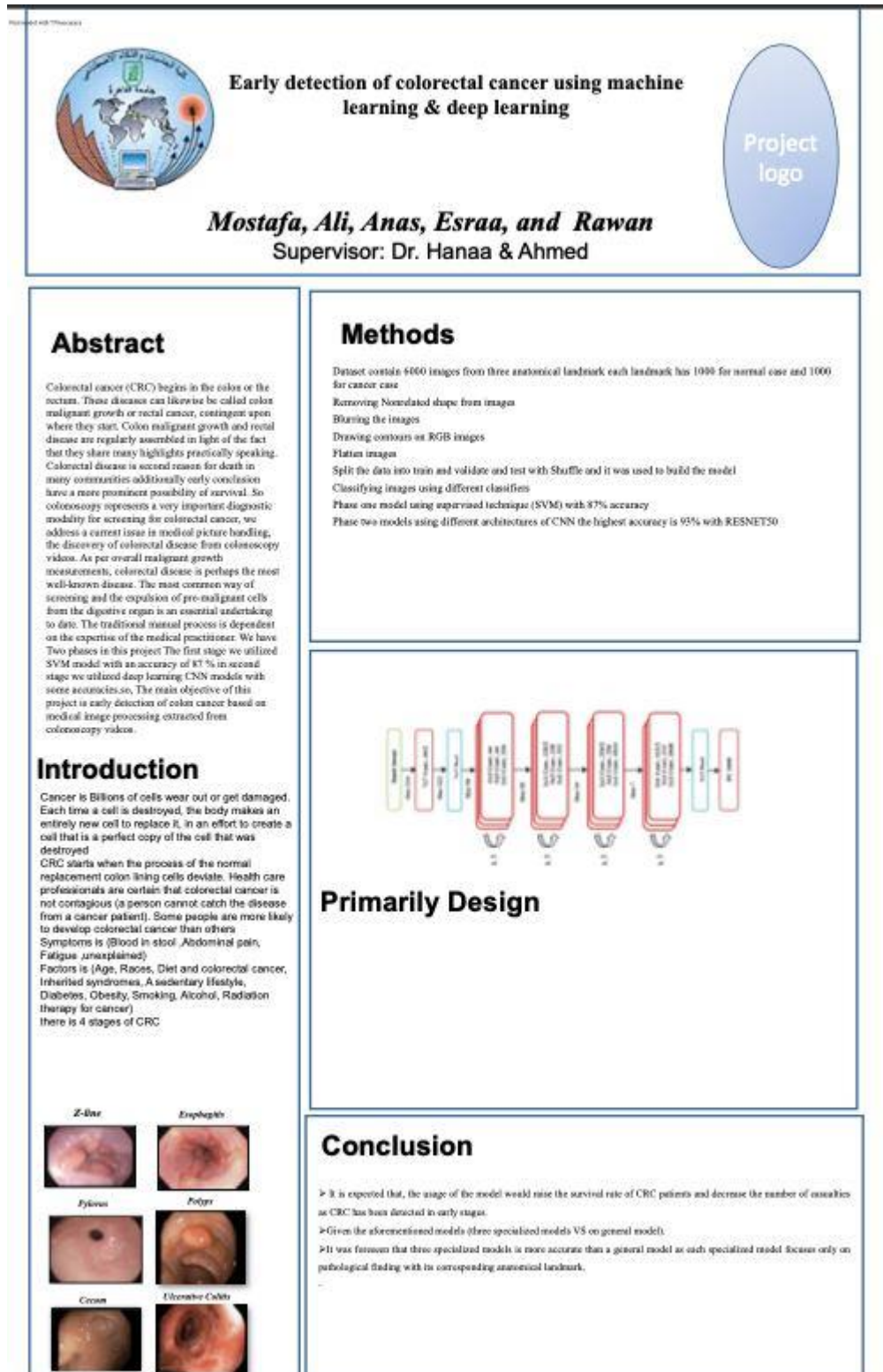
Fourth model CNN ResNet50 with 93% accuracy

```
inputs = Input(shape=(224,224,3))
base_model = ResNet50(include_top=False, weights='imagenet', input_shape=(224, 224,3))
base_model.trainable = False

added_layers = base_model(inputs, training=False)
added_layers = Flatten()(added_layers)
added_layers = Dense(100,activation='relu')(added_layers)
added_layers = Dense(50, activation='relu')(added_layers)
final_layer = Dense(6, activation='softmax')(added_layers)

KerasModel = Model(inputs, final_layer)
KerasModel.compile(optimizer='adam', loss='categorical_crossentropy', metrics=['accuracy'])
epochs = 25
KerasModel.fit(x_train, to_categorical(y_train), epochs=epochs, batch_size=64, verbose=1,
                validation_data=(x_valid, to_categorical(y_valid)))
```

Poster3



References

- Xu, L., Walker, B., Liang, P.-I., Tong, Y., Xu, C., Su, Y. C., & Karsan, A. (2020, August 21). Colorectal cancer detection based on Deep Learning. *Journal of pathology informatics*.
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- Sarwinda, D., Paradisa, R. H., Bustamam, A., & Anggia, P. (2021, February 19). Deep learning in image classification using residual network (ResNet) variants for detection of colorectal cancer. *Procedia Computer Science*.
- Abdullah M, Rani AA, Simadibrata M, Fauzi A, Syam AF (2012) The value of fecal tumor M2 pyruvate kinase as a diagnostic tool for colorectal cancer screening. *Acta med Indones* 44:94–99.