# **Project Report**

#### 1. INTRODUCTION

- 1.1 Project Overview
- 1.2 Purpose

#### 2. LITERATURE SURVEY

- 2.1 Existing problem
- 2.2 References
- 2.3 Problem Statement Definition

### 3. IDEATION & PROPOSED SOLUTION

- 3.1 Empathy Map Canvas
- 3.2 Ideation & Brainstorming
- 3.3 Proposed Solution
- 3.4 Problem Solution fit

## 4. REQUIREMENT ANALYSIS

- 4.1 Functional requirement
- 4.2 Non-Functional requirements

#### 5. PROJECT DESIGN

- 5.1 Data Flow Diagrams
- 5.2 Solution & Technical Architecture
- 5.3 User Stories

#### 6. PROJECT PLANNING & SCHEDULING

- 6.1 Sprint Planning & Estimation
- 6.2 Sprint Delivery Schedule
- 6.3 Reports from JIRA

## 7. CODING & SOLUTIONING (Explain the features added in the project along with code)

- 7.1 Feature 1
- 7.2 Feature 2
- 7.3 Database Schema (if Applicable)

#### 8. TESTING

- 8.1 Test Cases
- 8.2 User Acceptance Testing

#### 9. RESULTS

9.1 Performance Metrics

## 10. ADVANTAGES & DISADVANTAGES

- 11. CONCLUSION
- 12. FUTURE SCOPE

#### 13. APPENDIX

Source Code

GitHub & Project Demo Link

## INTRODUCTION

# 1.1. Project Overview

Now a day's people are suffering from skin diseases, more than 125 million people suffering from Psoriasis also skin cancer rate is rapidly increasing over the last few decades especially Melanoma is most diversifying skin cancer. If skin diseases are not treated at an earlier stage, then it may lead to complications in the body including spreading of the infection from one individual to the other. The skin diseases can be prevented by investigating the infected region at an early stage. The characteristic of the skin images is diversified so that it is a challenging job to devise an efficient and robust algorithm for automatic detection of skin disease and its severity. Skin tone and skin color play an important role in skin disease detection. Color and coarseness of skin are visually different. Automatic processing of such images for skin analysis requires quantitative discriminator to differentiate the diseases.

# 1.2. Project Purpose

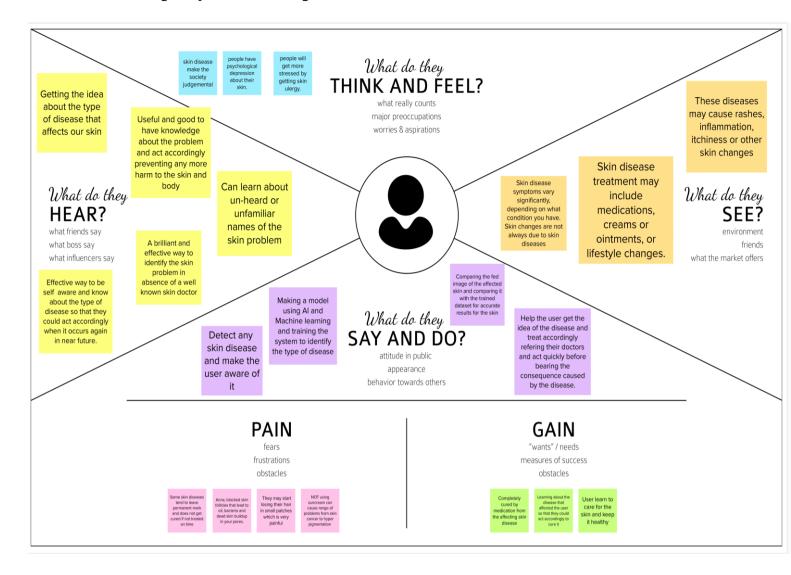
The goal is to identify and find cure for the affecting skin disease in an efficient and easy way possible by using advance artificial intelligence. We make it possible by building a model which is used for the prevention and early detection of skin cancer, psoriasis. Basically, skin disease diagnosis depends on the different characteristics like colour, shape, texture etc. Here the person can capture the images of skin and then the image will be sent the trained model. The model analyses the image and detect whether the person is having skin disease or not.

# LITERATURE SURVEY

- 2.1 Existing Problem
- 2.2 References
- 2.3 Problem Statement Definition

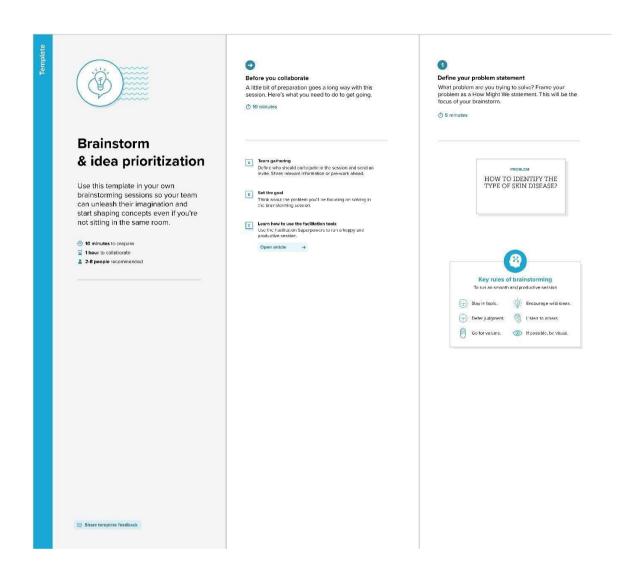
# IDEATHON AND PROPOSED SOLUTION

# 3.1 Empathy Canvas Map

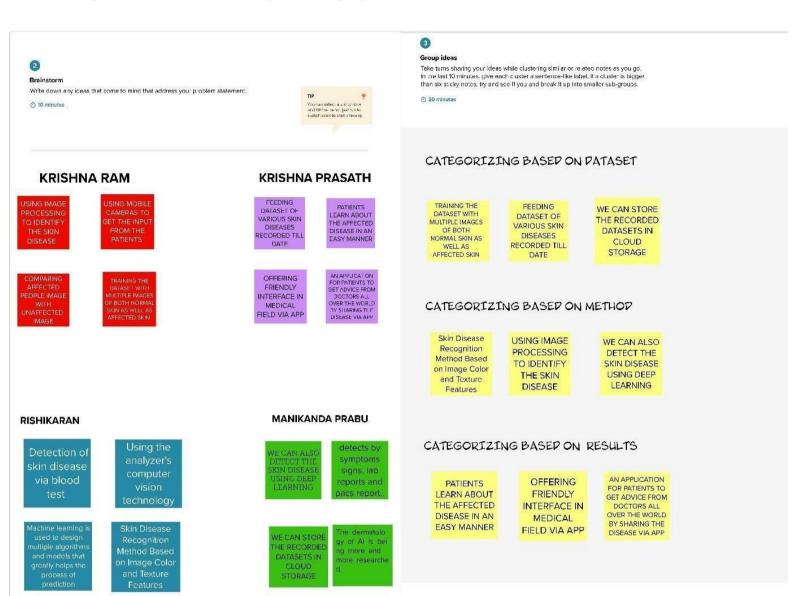


# 3.2 Ideation & Brainstorming

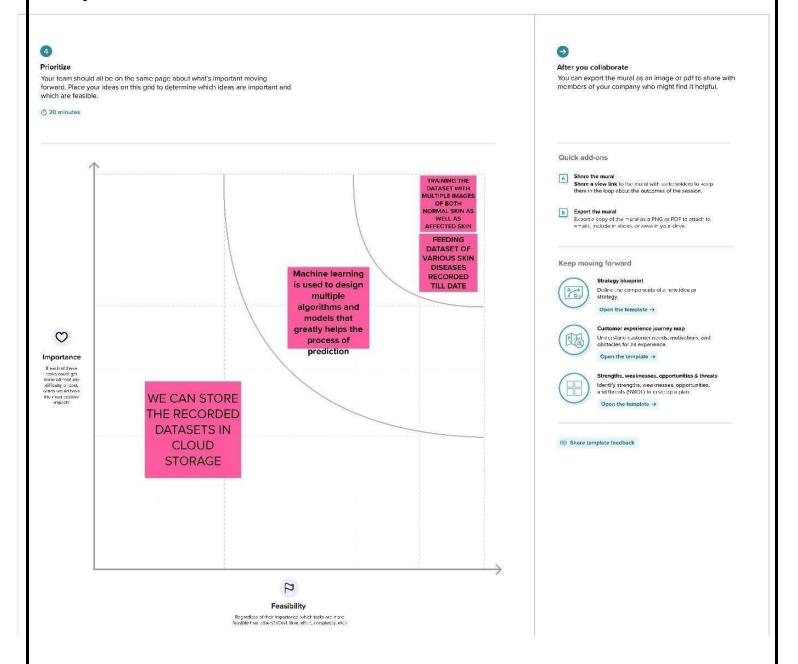
# Step-1: Team Gathering, Collaboration and Select the Problem Statement



### Step-2: Brainstorm, Idea Listing and Grouping



## **Step-3: Idea Prioritization**



# 3.3 Proposed Solution

S.No.	Parameter	Description
1.	Problem Statement (Problem to be solved)	Skin diseases vary in types, we are given the problem to identify the type of skin disease using AI based localization with Erythema. Our main goal is to identify the type of disease that the patient is affected with to make them aware and start to proceed by treating their skin.

2.	Idea / Solution description	Our idea is to train the machine with various
	Laster, Soldworf description	types of skin diseases by applying the
		knowledge of Data Science and Advance
		Machine Learning to make it capable for
		identifying the skin disease by using the set of fed Dataset. The machine uses advanced
		algorithms and image processing method to
		achieve the main goal/objective.
3.	Novelty / Uniqueness	The machine will be trained with the latest version of YOLO which currently is YOLOv5. Compared to other solutions performances, this machine is set to be fast and accurate in terms of mean average precision (mAP) and intersection over union (iOU) as well. It runs significantly faster than other detection methods with
		comparable performance.
		The images that are trained to follow the algorithm are trained by using Microsoft Visual Object Tagging Tool or VOTT in short, which is an open-source annotation and labelling tool for image and video assets including features such as; The ability to label images or video frames,
		extensible model for importing data from local or cloud storage providers, and model for exporting labelled data to local or cloud storage providers.
		For the current project, our datasets and image annotate will be stored in Cloudant DB, which is a non-relational, distributed database service provided by IBM.
4.	Social Impact / Customer Satisfaction	
		O With our product the patients will learn about the type of disease that had affected them so that they could act accordingly to cure it.
		O In the absence of a skin specialist, the machine can identify the affecting disease and hence the patient will be able to look for a cure rather than waiting for a doctor.
		O Providing a user-friendly interface and an ease for the patients to use it.
5.	Business Model (Revenue Model)	► First and foremost, benefit from this
		model is the reduction of time consumption for identification of any unknown skin disease.

#### 3.4 **Problem Solution fit**

Project Title: AI-based localization and classification of skin disease with erythema Project Design Phase-I - Solution Fit Template

CS

J&P

TR

**EM** 

Team ID: PNT2022TMID0010

Explore AS, differentiate

AS

# Define CS ≢ 00

### 1. CUSTOMER SEGMENT(S)

Who is your customer? i.e. working parents of 0-5 y.o. kids

Our customers are those people aged between 10-55 yr who are affected by severe skin diseases like erythema and skin cancer

#### 6. CUSTOMER CONSTRAINTS

constraints prevent your customers from taking action or limit their choices utlons? i.e. spending power, budget, no cash, network connection, available

The main constraint is financial support, as skin is important to be cared in regular time interval by skilled skin specialist.
second factor stands for limitations in their basic diet for curing the skin disease and maintaining the cycle for longer period of time. third factor is their time, the person goes through multiple diagnostics for learning about the skin disease which might takes a long time to be identified depending on the resources they use.

#### 5. AVAILABLE SOLUTIONS

CC

RC

SL

solutions are available to the customers when they face the to get the job done? What have they tried in the past? What pros & se solutions have? i.e. pen and paper is an altern

The most common and effective means of erythema detection is visual inspection of the skin. However, for people with darkly pigmented skin, erythema can be masked by melanin. Tissue Reflectance Spectroscopy (TRS) is a noninvasive method of quantifying skin color

됬

#### 2. JOBS-TO-BE-DONE / PROBLEMS

We help our customers to identify the problem/disease that they are affected with the help of modern technology such as AI/ML.

#### 9. PROBLEM ROOT CAUSE

What is the real reason that this problem exis What is the back story behind the need to do i.e. customers have to do it because of the change in regulations.

The major causes of the erythema are

- Herpes simplex virus Mycoplasma pneumoniae cytomegalovirus

#### 7. BEHAVIOUR

8. CHANNELS of BEHAVIOUR

What does your customer do to address the problem and get the job done? i.e. directly related: find the right solar panel installer, calculate usage and benefits; indirectly associated: customers spend free time on volunteering work (i.e.

At first, people start searching in internet for solutions and asking neighbors for medicines and tablets to cure by themselves in case home remedies don't work they reach out to the doctor and take the personal

What kind of actions do customers take online? Extract online channels from #7

What kind of actions do customers take offline? Extract offline channels from #7 and use them for customer development.

What triggers customers to act? i.e. seeing their neighbour installing solar panels, reading about a more efficient solution in the news

Basically people start to act or approach us when they find some unusuality in their day to day life for eg take a skin disease like erthyema. Erythema nodosum usually is caused by a reaction to a drug. an infection (bacterial, fungal, or viral), or another disorder such as inflammatory bowel disease. Typical symptoms include fever, joint pain, and characteristic painful red bumps and bruises on the person's shins and another example is skin cancer

Anyone who spends considerable time in the sun may develop skin cancer, especially if the skin isn't protected by sunscreen or clothing. So if our customers face these kind of problem they will start to take some

#### 4. EMOTIONS: BEFORE / AFTER

How do customers feel when they face a problem or a job and afterwards? i.e. lost, insecure > confident, in control - use it in your communication strategy & design.

After: Skin diseases have an adverse impact on psychosocial well-being and can lead to more depressive symptoms, social isolation, loneliness and decreased quality of life.

#### 10. YOUR SOLUTION

time consumption

skin specialist and

reduced to a millennial

If you are working on an existing busin s, write down your current solution first fill in the canvas, and check how much it fits reality.

If you are working on a new business proposition, then keep it blank until you fill in the canvas and come up with a solution that fits within customer limitations, solves a problem and matches customer behaviour.

First and foremost, benefit from this model is the reduction of

Treatment can be proceeded even without the presence of a

hence the chances for avoiding major effect on the skin is

for identification of any unknown skin disease.

8.2 OFFLINE

Their first move is to go and look up for the symptoms and effects of the disease that affected them. Looking for a home remedy solution from

They physically reach out to a skin specialist to gain knowledge about th skin disease and take remedy from them first hand.

#### Online:

network or online skin specialist blogpost etc.

СН

ify strong TR & EM



# REQUIREMENT ANALYSIS

# **4.1** Functional requirements

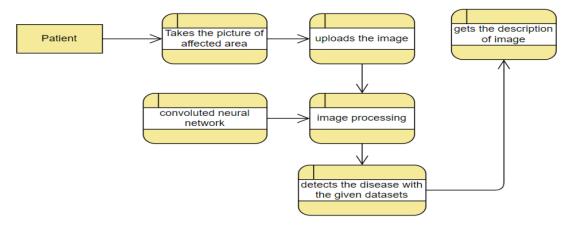
FR No.	Functional Requirement (Epic)	Sub Requirement (Story / Sub-Task)
FR-1	User Registration	Registration through Form
		Registration through Gmail
FR-2	User Confirmation	Confirmation via Email
		Confirmation via OTP
FR-3	User Profile	Users provides their medical history.
FR-4	User Uploads Images (Input)	Upload the images as jpeg/jpg's Upload images as png's
FR-5	Output Analysis	Output analyzed through trained model
FR-6	Provides Description	Gives the detailed description of the skin disease found

# **4.2** Non-Functional requirements

FR No.	Non-Functional Requirement	Description
NFR-1	Usability	Used to classify skin disease with erythema
NFR-2	Security	It offers greater security and prevents unauthorized individuals from accessing user's data.
NFR-3	Reliability	Even with more users, there will be a good performance without failure.
NFR-4	Performance	With greater accuracy, the performance is high.
NFR-5	Availability	With a good system, all authorized users can access it.
NFR-6	Scalability	Performance will be good even with the higher user traffic,

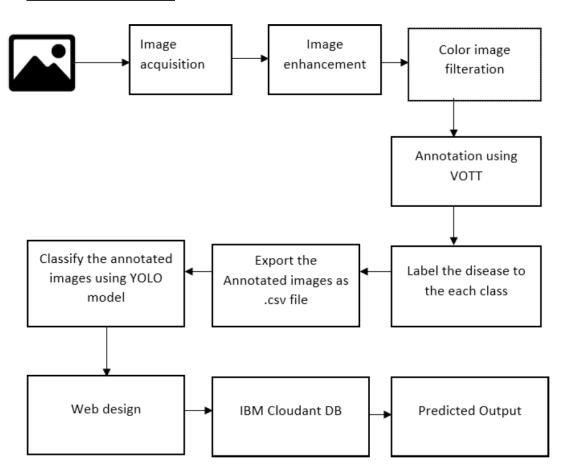
# **PROJECT DESIGN**

# **5.1** Data Flow Diagram

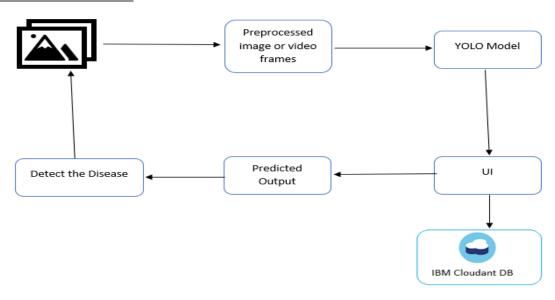


# 5.2 Solution & Technical Architecture

Solution Architecture:-



# Technical Architecture:-



# 5.3 User Stories

User Type	Functional Requirement (Epic)	User Story Number	User Story / Task	Acceptance criteria	Priority	Release
Customer (Mobile user)	Registration	USN-1	As a user, I can register for the application by entering my email, password, and confirming my password.	I can access my account / dashboard	High	Sprint-1
	Confirm	USN-2	As a user, I will receive confirmation email once I have registered for the application	I can receive confirmation email & click confirm	High	Sprint-1
	Registration	USN-3	As a user, I can register for the application through Facebook	I can register & access the dashboard with Facebook Login	High	Sprint-2
	Registration	USN-4	As a user, I can register for the application through Gmail	I can access my account / dashboard	Medium	Sprint-1
	Login	USN-5	As a user, I can log into the application by entering email & password	I can access Dashboard	High	Sprint-1
	Dashboard	USN-6	As a user, I can Access my Dashboard.	I can interact with the inteface	Medium	Sprint-3
	Data Input	USN-7	As a user, I can upload the images of the affected skin area	I can submit it to the application	High	Sprint-4
Customer Care	Solution	USN-8	Responding to each email you receive	Offer a solution for how your	High	Sprint-3

User Type	Functional Requirement (Epic)	User Story Number	User Story / Task	Acceptance criteria	Priority	Release
Executive				company can improve the customer's experience.		
Administrator	Manage	USN-9	Do-it-yourself service for delivering Everything.	set of predefined requirements that must be met to mark a user story complete.	High	Sprint-4
Training Model	Image processing	USN-10	By comparing the images, the disease will be detected with the given datasets	All the necessary operation performed and information extracted	High	Sprint-3
	Report generation	USN-11	Based on the detection of disease, report generated	The results will be shown on the screen to the patients	High	Sprint-4

# PROJECT PLANNING & SCHEDULING

# **6.1** Sprint Planning & Estimation

Sprint	Functional Requirement (Epic)	User Story Number	User Story / Task	Story Points	Priority	Team Members
Splint-1	Pre-requisites	USN-1	Install Python IDE, Python packages, Microsoft Visual Object Tagging Tool, Yolo Structure	7	High	Krishna Manikanda Krishna prasath
Sprint-1	Data Collection	USN-2	The dataset should be collected in realtime or from the gallery or collect it from google.	10	High	Krishna Manikanda
Splint-1	Annotate Images	USN-3	Create a project in Visual Object Tagging Tool	3	Medium	Rishikaran Krishna prasath
Splint-2	Training YOLO	USN-4	In this we will train our model using YOLO weights	5	Medium	Krishna Rishikaran
Splint-2		USN-5	Download and convert pre-trained weights	5	High	Manikanda Rishikaran
Splint-2		USN-6	To start training run the training script within the YOLO structure.	10	Low	Krishna manikanda
Splint-3	Cloudant DB	USN-7	Register and Login to IBM Cloud	5	Medium	Rishikaran Krishna prasath
Sprint-3		USN-8	Create Service Instant and credentials	5	High	Rishikaran Krishna prasath

Sprint	Functional Requirement (Epic)	User Story Number	User Story / Task	Story Points	Priority	Team Members
Sprint-3		USN-9	Launch Cloudant DB and then create database	2	High	manikanda Krishna prasath
Sprint-3	Developing Phase	USN-10	In this build a web application that is integrated to the caffemodel.	3	Low	Krishna Rishikaran
Sprint-3		USN-11	For this build HTML Pages	2	Medium	manikanda Krishna prasath
Sprint-3		USN-12	Develop and build the python code to run the application.	3	High	Rishikaran Krishna prasath
Sprint-4	Testing Phase	USN-13	As a user login to the dashboard	10	High	Krishna manikanda
Sprint-4		USN-14	As a user import the skin affected disease image to the software application.	5	Medium	Rishikaran Krishna prasath
Sprint-4		USN-15	YOLO will process the image and give the result as unaffected or affected with other details	5	Medium	Krishna Rishikaran

# **6.2** Sprint Delivery Schedule

S.no	Milestone	Activities	Start Date	End Date
1	Project Objectives	Prepare the project objectives.	22-Aug-2022	27-Aug-2022
2	Pre-Requisites	<ul> <li>Install Python IDE</li> <li>Install Microsoft's Visual Object Tagging Tool</li> <li>Download YOLO Project Structure</li> </ul>	22-Aug-2022	27-Aug-2022
3	Create Dataset	Creating Datasetfromscratch.	22-Aug-2022	27-Aug-2022
4	Annotate Images	Create a project in VOTT  VOTT Project Creation.	27-Aug-2022	02-Sep-2022
5	Training YOLO	Download and Convert Pre-Trained weights. Train YOLOV3 Detector	24-oct-2022	19-Nov-2022
6	Cloudant DB	<ul> <li>Register and login to the IBM cloud</li> <li>Create service instance</li> <li>Create service credentials.</li> <li>Launch Cloudant DB</li> <li>Create Database</li> </ul>	24-oct-2022	12-Nov-2022
7	Application Building	<ul> <li>➢ Build HTML page</li> <li>➢ Build PYTHON code</li> <li>➢ Run the application</li> </ul>	24-oct-2022	19-Nov-2022
8	Ideation Phase	<ul> <li>➤ Literature Survey</li> <li>➤ Empathy Map</li> <li>➤ Ideation</li> </ul>	29-Aug-2022	01-Oct -2022

.no	Milestone	Activities	Start Date	End Date
	Project Design Phase - I	<ul> <li>Proposed Solution</li> <li>Problem Solution Fit</li> <li>Solution Architecture</li> </ul>	19-Sept-2022	01-Oct-2022
0	Project Design Phase - II	<ul> <li>Customer Journey</li> <li>Functional Requirement</li> <li>Data Flow Diagrams</li> <li>Technology Architecture</li> </ul>	03-Oct-2022	15-Oct-2022
1	Project Planning Phase	<ul> <li>Prepare Milestone &amp; Activity List</li> <li>Sprint Delivery Plan</li> </ul>	17-Oct-2022	21-Oct-2022
2	Project Development Phase	<ul> <li>Project Development - Delivery of Sprint-1</li> <li>Project Development - Delivery of Sprint-2</li> <li>Project Development - Delivery of Sprint-3</li> <li>Project Development - Delivery of Sprint-4</li> </ul>	24-Oct-2022	19-Nov-2022

## CODING AND SOLUTIONING

```
7.1
      Feature 1
//Index Page HTML
<!DOCTYPE html>
<html lang="en">
<head>
  <meta charset="UTF-8">
  <meta http-equiv="X-UA-Compatible" content="IE=edge">
  <meta name="viewport" content="width=device-width, initial-scale=1.0">
  rel="stylesheet"
                           href="https://maxcdn.bootstrapcdn.com/bootstrap/4.0.0/css/bootstrap.min.css"
integrity="sha384-
Gn5384xqQ1aoWXA+058RXPxPg6fy4IWvTNh0E263XmFcJlSAwiGgFAW/dAiS6JXm"
crossorigin="anonymous">
                    src="https://code.jquery.com/jquery-3.2.1.slim.min.js"
                                                                                  integrity="sha384-
KJ3o2DKtlkvYIK3UENzmM7KCkRr/rE9/Qpg6aAZGJwFDMVNA/GpGFF93hXpG5KkN"
crossorigin="anonymous"></script>
                          src="https://cdnjs.cloudflare.com/ajax/libs/popper.js/1.12.9/umd/popper.min.js"
integrity="sha384-ApNbgh9B+Y1QKtv3Rn7W3mgPxhU9K/ScQsAP7hUibX39j7fakFPskvXusvfa0b4Q"
crossorigin="anonymous"></script>
  <script src="https://maxcdn.bootstrapcdn.com/bootstrap/4.0.0/js/bootstrap.min.js"</pre>
                                                                                 integrity="sha384-
JZR6Spejh4U02d8jOt6vLEHfe/JQGiRRSQQxSfFWpi1MquVdAyjUar5+76PVCmYl"
crossorigin="anonymous"></script>
  <script src="https://kit.fontawesome.com/8b9cdc2059.js" crossorigin="anonymous"></script>
                 href="https://fonts.googleapis.com/css2?family=Akronim&family=Times"
                                                                                              New
Roman&display=swap" rel="stylesheet">
  <title>SKINBOSS</title>
</head>
<body>
<style>
.icon-bar {
 width: 90px; /* Set a specific width */
 background-color: #555; /* Dark-grey background */
.icon-bar a {
 display: block; /* Make the links appear below each other instead of side-by-side */
 text-align: center; /* Center-align text */
 padding: 16px; /* Add some padding */
 transition: all 0.3s ease; /* Add transition for hover effects */
 color: white; /* White text color */
 font-size: 36px; /* Increased font-size */
```

```
.icon-bar a:hover {
 background-color:black; /* Add a hover color */
.active {
 background-color:grey; /* Add an active/current color */
.nav--items {
 overflow: hidden;
 background-color: #f1f1f1;
/* Style the buttons that are used to open the tab content */
.nav--items a {
 color:black;
 background-color: inherit;
 float: right;
 border: none;
 outline: none:
 cursor: pointer;
 padding: 14px 16px;
 transition: 0.3s;
/* Change background color of buttons on hover */
.nav--items a:hover {
 background-color: #ddd;
.nav--items a.active {
       color:black;
 background-color: #ccc;
.heading{
text-align:center;
color:white;
background-color:#22DDCA;
.top{
background-color:#7BEADF;
p{
color:black;
.head{
font-family:timesnewroman;
color:black;
text-align:center;
```

```
}
</style>
  <header id="head" class="header">
 <section id="navbar">
      <h1 class="heading">SKIN BOSS</h1>
   <div class="nav--items">
      <a href="{{url_for('login')}}">Log In</a>
      <a href="{{url_for('signup')}}">Sign Up</a>
        <a href="{{url_for('logout')}}">Log Out</a>
             <a href="{{url_for('prediction')}}">Prediction</a>
   </div>
 </section>
 <div class="top">
  <h2 class="title text-muted">
    A PERFECT LIFE WITH PERFECT SKIN
  </h2>
</div>
 <section id="slider">
  <div id="carouselExampleIndicators" class="carousel" data-ride="carousel">

    class="carousel-indicators">

      data-target="#carouselExampleIndicators" data-slide-to="0" class="active ">
      data-target="#carouselExampleIndicators" data-slide-to="2">
    </01>
    <div class="carousel-item active">
                     class="d-block
                                             w-100"
                                                              src="https://regencyhealthcare.in/wp-
content/uploads/2021/04/Artboard-5-1.png" alt="First slide" style="height:500px">
    </div>
    <div class="carousel-item">
      <img alt="Second slide" class="d-block w-100" src="https://img.graphicsurf.com/2020/10/skin-</pre>
disease-vector-design-concept.jpg" style="height:500px">
    </div>
    <a class="carousel-control-prev" href="#carouselExampleIndicators" role="button" data-slide="prev">
      <span class="carousel-control-prev-icon" aria-hidden="true"></span>
      <span class="sr-only">Previous</span>
    </a>
    <a class="carousel-control-next" href="#carouselExampleIndicators" role="button" data-slide="next">
      <span class="carousel-control-next-icon" aria-hidden="true"></span>
      <span class="sr-only">Next</span>
    </a>
  </div>
   <div class="Problem">
```

```
<h1 class="head">Problem Statement</h1>
```

Now a day's people are suffering from skin diseases, More than 125 million people suffering from Psoriasis also skin cancer rate is rapidly increasing over the last few decades especially Melanoma is most diversifying skin cancer. If skin diseases are not treated at an earlier stage, then it may lead to complications in the body including spreading of the infection from one individual to the other. The skin diseases can be prevented by investigating the infected region at an early stage. The characteristic of the skin images is diversified so that it is a challenging job to devise an efficient and robust algorithm for automatic detection of skin disease and its severity. Skin tone and skin colour play an important role in skin disease detection. Colour and coarseness of skin are visually different. Automatic processing of such images for skin analysis requires quantitative discriminator to differentiate the diseases.

To overcome the above problem we are building a model which is used for the prevention and early detection of skin cancer, psoriasis. Basically, skin disease diagnosis depends on the different characteristics like colour, shape, texture etc. Here the person can capture the images of skin and then the image will be sent the trained model. The model analyses the image and detect whether the person is having skin disease or not.

```
<h1 class="head">Proposed Solution</h1>

    Different skin disorders can be detected by just submitting photographs, and this approach is quite effective at helping people in the community identify ailments earlier.
    Our return on investment will be the creation and distribution of a proprietary product that will be used as a solution.
    This system is more scalable because it accepts any picture type, regardless of resolution, and offers good performance in any situation.

</header>

</body>
```

# 7.2 Feature 2 //Model to train

```
# -*- coding: utf-8 -*-
"""Untitled0.ipynb
Automatically generated by Colaboratory.
Original file is located at
https://colab.research.google.com/drive/1PYFZ7zKhWpFF5YilnguhZ8X1EgtSIJN4
"""
```

import re

</html>

```
import numpy as np
import os
from flask import Flask, app,request,render_template
import sys
from flask import Flask, request, render_template, redirect, url_for
import argparse
from tensorflow import keras
from PIL import Image
from timeit import default_timer as timer
import test
from pyngrok import ngrok
import pandas as pd
import numpy as np
import random
def get_parent_dir(n=1):
""" returns the n-th parent dicrectory of the current
working directory """
current_path = os.path.dirname(os.path.abspath(_file_))
for k in range(n):
current_path = os.path.dirname(current_path)
return current_path
src path=r'/content/drive/MyDrive/IBM PROJECT/yolo structure/2 Training/src'
print(src_path)
utils_path=r'/content/drive/MyDrive/IBM_PROJECT/yolo_structure/Utils'
print(utils_path)
sys.path.append(src_path)
sys.path.append(utils_path)
import argparse
from keras_yolo3.yolo import YOLO, detect_video
from PIL import Image
from timeit import default_timer as timer
from utils import load extractor model, load features, parse input, detect object
import test
import utils
import pandas as pd
import numpy as np
from Get_File_Paths import GetFileList
import random
os.environ["TF CPP MIN LOG LEVEL"] = "3"
# Set up folder names for default values
data_folder = os.path.join(get_parent_dir(n=1), "yolo_structure", "Data")
image folder = os.path.join(data folder, "Source Images")
image_test_folder = os.path.join(image_folder, "Test_Images")
detection results folder = os.path.join(image folder, "Test Image Detection Results")
detection results file = os.path.join(detection results folder, "Detection Results.csv")
model_folder = os.path.join(data_folder, "Model_Weights")
model weights = os.path.join(model folder, "trained weights final.h5")
model_classes = os.path.join(model_folder, "data_classes.txt")
anchors path = os.path.join(src path, "keras yolo3", "model data", "yolo anchors.txt")
```

```
FLAGS = None
from cloudant.client import Cloudant
# Authenticate using an IAM API key
client =
Cloudant.iam('ef7f4729-2486-45c5-a7fa-f4140373e2e6-bluemix','6GfFjs3engXLnSJB8Kp4f
bs7HTKwrJpWJE7wNPGzZPVW', connect=True)
# Create a database using an initialized client
my_database = client.create_database('my_database')
app=Flask( name )
port no=5000
ngrok.set_auth_token("2H7aM94zEuTa40t3J6jKpIqWAc3_B2UxzZs6qxetntgadxQW")
public url = ngrok.connect(port no).public url
print(f"To acces the Gloable link please click {public_url}")
#default home page or route
@app.route('/')
def index():
return render_template('index.html')
@app.route('/index.html')
def home():
return render template("index.html")
#registration page
@app.route('/register')
def register():
return render_template('register.html')
@app.route('/afterreg', methods=['POST'])
def afterreg():
x = [x \text{ for } x \text{ in request.form.values}()]
print(x)
data = {
'_id': x[1], # Setting _id is optional
'name': x[0],
'psw':x[2]
print(data)
query = {'_id': {'$eq': data['_id']}}
docs = my_database.get_query_result(query)
print(docs)
print(len(docs.all()))
if(len(docs.all())==0):
url = my database.create document(data)
#response = requests.get(url)
return render_template('register.html', pred="Registration Successful, please
login using your details")
else:
return render template('register.html', pred="You are already a member, please
```

```
login using your details")
#login page
@app.route('/login')
def login():
return render_template('login.html')
@app.route('/afterlogin',methods=['POST'])
def afterlogin():
user = request.form['_id']
passw = request.form['psw']
print(user,passw)
query = {'_id': {'$eq': user}}
docs = my_database.get_query_result(query)
print(docs)
print(len(docs.all()))
if(len(docs.all())==0):
return render_template('login.html', pred="The username is not found.")
if((user==docs[0][0]['\_id'] and passw==docs[0][0]['psw'])):
return redirect(url_for('prediction'))
else:
print('Invalid User')
@app.route('/logout')
def logout():
return render_template('logout.html')
@app.route('/prediction')
def prediction():
return render_template('prediction.html',path="../static/img/6623.jpg",)
@app.route('/result',methods=["GET","POST"])
def res():
# Delete all default flags
parser = argparse.ArgumentParser(argument_default=argparse.SUPPRESS)
Command line options
f = request.files['file']
f.save("./drive/MyDrive/IBM_PROJECT/Flask/static/img/"+f.filename)
parser.add_argument(
"--input_path",
type=str,
default=image_test_folder,
```

```
help="Path to image/video directory. All subdirectories will be included. Default
is "
+ image_test_folder,
parser.add_argument(
"--output",
type=str,
default=detection_results_folder,
help="Output path for detection results. Default is "
+ detection results folder,
parser.add_argument(
"--no_save_img",
default=False,
action="store_true",
help="Only save bounding box coordinates but do not save output images with
annotated boxes. Default is False.",
parser.add_argument(
"--file_types",
"--names-list",
nargs="*",
default=[],
help="Specify list of file types to include. Default is --file_types .jpg .jpeg .png
.mp4",
parser.add_argument(
"--yolo_model",
type=str,
dest="model_path",
default=model_weights,
help="Path to pre-trained weight files. Default is " + model_weights,
parser.add_argument(
"--anchors",
type=str,
dest="anchors_path",
default=anchors_path,
help="Path to YOLO anchors. Default is " + anchors path,
parser.add_argument(
"--classes",
type=str,
dest="classes_path",
default=model classes,
help="Path to YOLO class specifications. Default is " + model_classes,
parser.add_argument(
"--gpu num", type=int, default=1, help="Number of GPU to use. Default is 1"
```

```
)
parser.add argument(
"--confidence",
type=float,
dest="score",
default=0.25,
help="Threshold for YOLO object confidence score to show predictions. Default is
0.25.",
parser.add_argument(
"--box_file",
type=str,
dest="box",
default=detection_results_file,
help="File to save bounding box results to. Default is "
+ detection_results_file,
parser.add_argument(
"--postfix",
type=str,
dest="postfix",
default="_disease",
help='Specify the postfix for images with bounding boxes. Default is "_disease",
yolo = YOLO(
**{
"model_path": FLAGS.model_path,
"anchors_path": FLAGS.anchors_path,
"classes_path": FLAGS.classes_path,
"score": FLAGS.score,
"gpu_num": FLAGS.gpu_num,
"model_image_size": (416, 416),
img_path="/drive/MyDrive/IBM_PROJECT/Flask/static/img/"+f.filename
prediction, image,lat,lon= detect_object(
yolo,
img_path,
save img=save img,
save_img_path=FLAGS.output,
postfix=FLAGS.postfix,
yolo.close_session()
return
render_template('prediction.html',prediction=str(prediction),path="../static/img/"+f.filena
me)
```

```
""" Running our application """ if _name_ == "_main_": app.run(port=port_no)
```

## 7.3 Database Schema

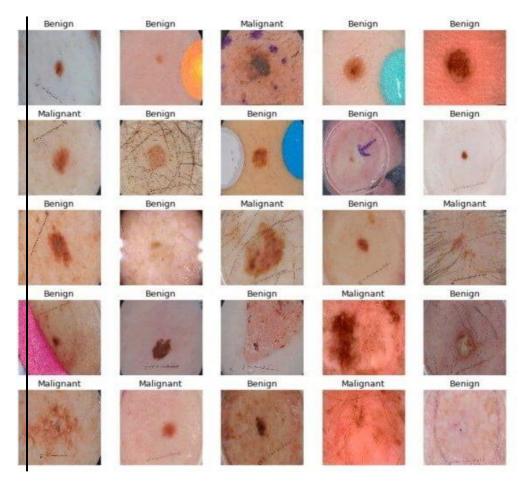
Currently we will be using the cloudantDB provided by IBM for storing all annotated images for already identified skin diseases and store in the cloud sever. As the images are scanned, the similar images are pulled out from the server for identifing the disease nature.

#### Output:

Number of training samples: 2000 Number of validation samples: 150

```
# preprocess data def
    decode_img(img):
    # convert the compressed string to a 3D uint8 tensor
     img = tf.image.decode_jpeg(img, channels=3)
 # Use `convert_image_dtype` to convert to floats in the [0,1] range.img
     = tf.image.convert_image_dtype(img, tf.float32)
 # resize the image to the desired size.
     return tf.image.resize(img, [299, 299])
def process path(filepath, label):
 # load the raw data from the file as a stringing
     = tf.io.read_file(filepath)
 img = decode_img(img)
    return img, label
valid_ds = valid_ds.map(process_path)
    train_ds = train_ds.map(process_path)#
    test_ds = test_ds
    for image, label in train_ds.take(1):
      print("Image shape:", image.shape)
      print("Label:", label.numpy())
Image shape: (299, 299, 3)
Label: 0
# training parameters
    batch size = 64
    optimizer = "rmsprop"
```

```
def prepare_for_training(ds, cache=True, batch_size=64, shuffle_buffer_size=1000):if
     cache:
  if isinstance(cache, str):
       ds = ds.cache(cache)
      else:
   ds = ds.cache()
 # shuffle the dataset
 ds = ds.shuffle(buffer_size=shuffle_buffer_size)#
     Repeat forever
 ds = ds.repeat() #
     split to batches
 ds = ds.batch(batch\_size)
 #`prefetch` lets the dataset fetch batches in the background while the model# is
     training.
 ds = ds.prefetch(buffer_size=tf.data.experimental.AUTOTUNE)return
     ds
valid_ds = prepare_for_training(valid_ds, batch_size=batch_size, cache="valid-cached-data")
    train_ds = prepare_for_training(train_ds, batch_size=batch_size, cache="train-cached-data")
    batch = next(iter(valid_ds))
    def show_batch(batch):
     plt.figure(figsize=(12,12))
     for n in range (25):
   ax = plt.subplot(5,5,n+1)
       plt.imshow(batch[0][n])
   plt.title(class_names[batch[1][n].numpy()].title()) plt.axis('off')
show_batch(batch)
    Output:
```



```
# building the model
# InceptionV3 model & pre-trained weights
module_url = "https://tfhub.dev/google/tf2-preview/inception_v3/feature_vector/4"

m = tf.keras.Sequential([
          hub.KerasLayer(module_url, output_shape=[2048], trainable=False),
          tf.keras.layers.Dense(1, activation="sigmoid")
])

m.build([None, 299, 299, 3])
m.compile(loss="binary_crossentropy", optimizer=optimizer, metrics=["accuracy"])m.summary()
```

# Output:

## Model: "sequential"

Layer (type)	Output Shape	Param #	
keras_layer (Keras	Layer) multiple	21802784	
dense (Dense)	multiple	2049	=========

Total params: 21,804,833 Trainable params: 2,049

Non-trainable params: 21,802,784

\_\_\_\_

## Training the Model

## Output:

```
Train for 31 steps, validate for 2 steps
 Epoch 1/100
 Epoch 00001: val_loss improved from inf to 0.49703, saving model to benign-vs-
 malignant_64_rmsprop_0.497.h5
0.4970 - val_accuracy: 0.8125
<..SNIPED..>
Epoch 27/100
Epoch 00027: val_loss improved from 0.40253 to 0.38991, saving model to benign-vs-
 malignant_64_rmsprop_0.390.h5
val_loss: 0.3899 - val_accuracy: 0.8359
<..SNIPED..>
Epoch 41/100
Epoch 00041: val_loss did not improve from 0.38991
val_loss: 0.3948 - val_accuracy: 0.8281
 Epoch 42/100
Epoch 00042: val_loss did not improve from 0.38991
val_loss: 0.4572 - val_accuracy: 0.8047
```

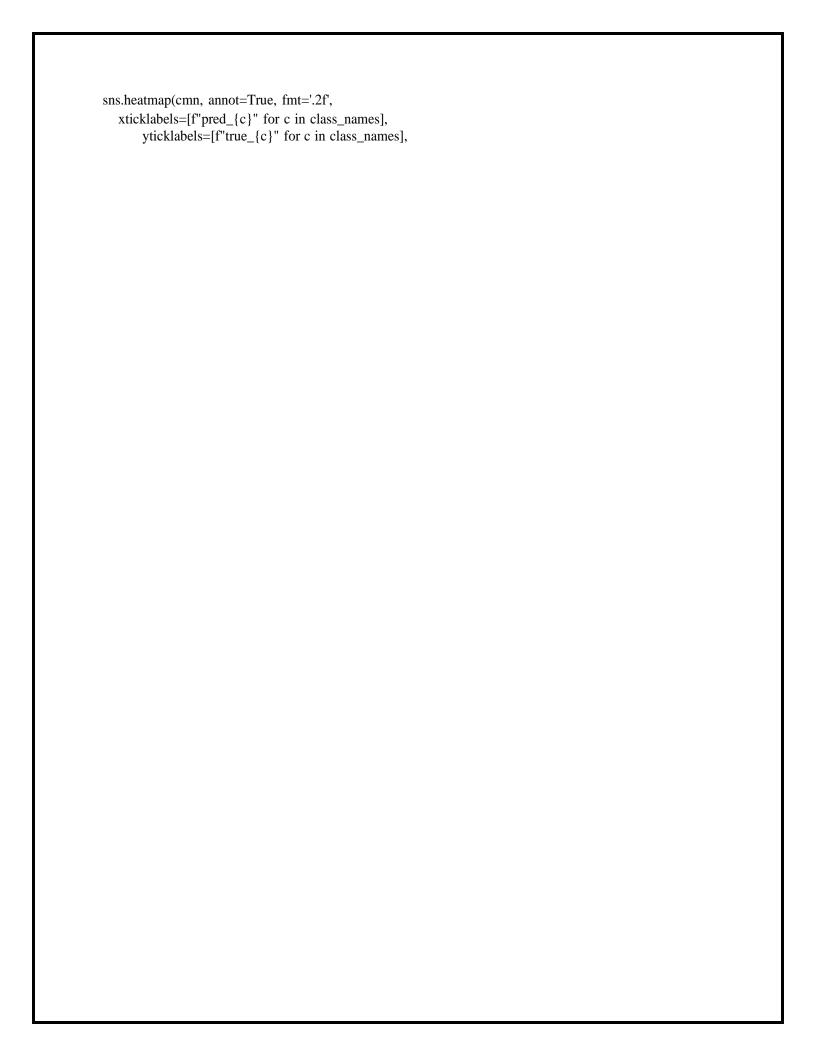
#### **Model Evaluation:**

```
# load testing set test_metadata_filename
    = "test.csv"
    df_test = pd.read_csv(test_metadata_filename)
    n_testing_samples = len(df_test)
    print("Number of testing samples:", n_testing_samples)
    test_ds = tf.data.Dataset.from_tensor_slices((df_test["filepath"], df_test["label"]))def
    prepare_for_testing(ds, cache=True, shuffle_buffer_size=1000):
     if cache:
      if isinstance(cache, str):
       ds = ds.cache(cache)
      else:
       ds = ds.cache()
     ds = ds.shuffle(buffer_size=shuffle_buffer_size)
     return ds
    test_ds = test_ds.map(process_path)
    test_ds = prepare_for_testing(test_ds, cache="test-cached-data")
Number of testing samples: 600#
    evaluation
# load testing set test_metadata_filename
    = "test.csv"
df_test = pd.read_csv(test_metadata_filename)
    n_testing_samples = len(df_test)
print("Number of testing samples:", n_testing_samples)
test ds = tf.data.Dataset.from tensor slices((df test["filepath"], df test["label"]))
    def prepare_for_testing(ds, cache=True, shuffle_buffer_size=1000):if
     cache:
  if isinstance(cache, str):ds =
```

# evaluation

ds.cache(cache) else:			
ds = ds.cache()			

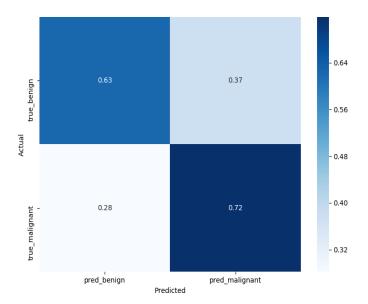
```
ds = ds.shuffle(buffer size=shuffle buffer size)return
test_ds = test_ds.map(process_path)
test_ds = prepare_for_testing(test_ds, cache="test-cached-data")
    # load the weights with the least loss
    m.load_weights("benign-vs-malignant_64_rmsprop_0.390.h5")
    print("Evaluating the model...")
    loss, accuracy = m.evaluate(X_test, y_test, verbose=0)
    print("Loss:", loss, " Accuracy:", accuracy)
    Output:
    Evaluating the model...
    Loss: 0.4476394319534302 Accuracy: 0.8
def get_predictions(threshold=None):
 Returns predictions for binary classification given `threshold`
 For instance, if threshold is 0.3, then it'll output 1 (malignant) for that sample if the
     probability of 1 is 30% or more (instead of 50%)
 y_pred = m.predict(X_test)if
     not threshold:
  threshold = 0.5
 result = np.zeros((n_testing_samples,))for i
     in range(n_testing_samples):
  # test melanoma probabilityif
      y_pred[i][0] >= threshold:
   result[i] = 1
      # else, it's 0 (benign)
     return result
threshold = 0.23
# get predictions with 23% threshold
# which means if the model is 23% sure or more that is malignant,# it's
    assigned as malignant, otherwise it's benign
y_pred = get_predictions(threshold)
def plot_confusion_matrix(y_test, y_pred):
 cmn = confusion_matrix(y_test, y_pred)#
    Normalise
 cmn = cmn.astype('float') / cmn.sum(axis=1)[:, np.newaxis]#
     print it
 print(cmn)
 fig, ax = plt.subplots(figsize=(10,10))
```



```
cmap="Blues"
)
plt.ylabel('Actual')
plt.xlabel('Predicted')
# plot the resulting confusion matrix
plt.show()

plot_confusion_matrix(y_test, y_pred)
```

# Output:



```
sensitivity = sensitivity_score(y_test, y_pred)
    specificity = specificity_score(y_test, y_pred)
print("Melanoma Sensitivity:", sensitivity)
    print("Melanoma Specificity:", specificity)
```

# Output:

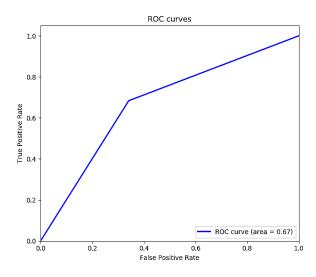
# Melanoma Sensitivity: 0.717948717948718

Melanoma Specificity: 0.6252587991718427

```
def plot_roc_auc(y_true, y_pred):
    """
    This function plots the ROC curves and provides the scores."""
        # prepare for figure
        plt.figure()
    fpr, tpr, _ = roc_curve(y_true, y_pred)
```

```
# obtain ROC AUC
      roc_auc = auc(fpr, tpr)
      # print score
      print(f"ROC AUC:
      {roc_auc:.3f}")# plot ROC
      curve
  plt.plot(fpr, tpr, color="blue", lw=2,
             label='ROC curve (area = {f:.2f})'.format(d=1, f=roc_auc))
      plt.xlim([0.0, 1.0])
  plt.ylim([0.0, 1.05])
      plt.xlabel('False Positive Rate')
      plt.ylabel('True Positive Rate')
      plt.title('ROC curves')
      plt.legend(loc="lower right")
      plt.show()
plot_roc_auc(y_test, y_pred)
```

# Output:

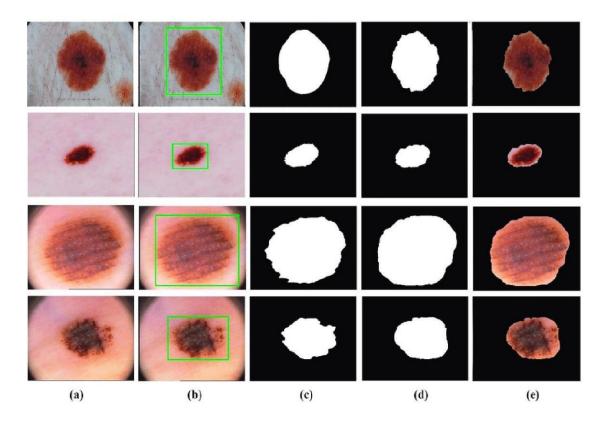


**ROC AUC: 0.671** 

# 8. TESTING

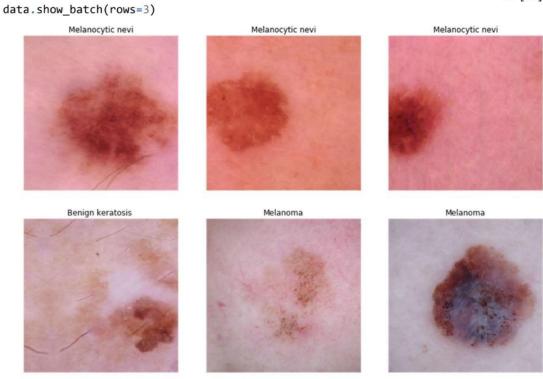
# **8.1 Test Cases**

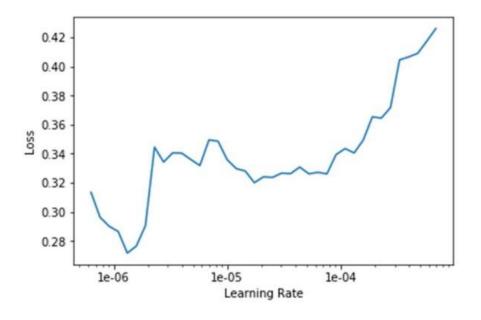
The final results are based on the accuracy results in the form of the melanoma and the non-melanomaskin diseases classifications.



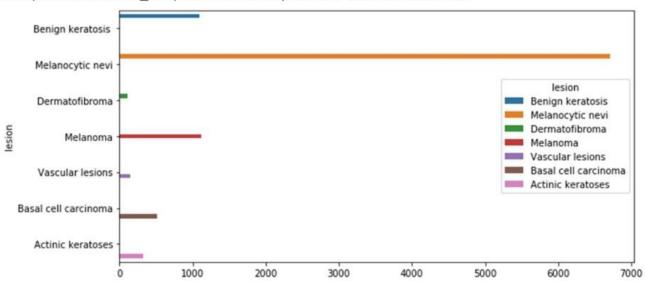
# 8.2 User Acceptance Testing



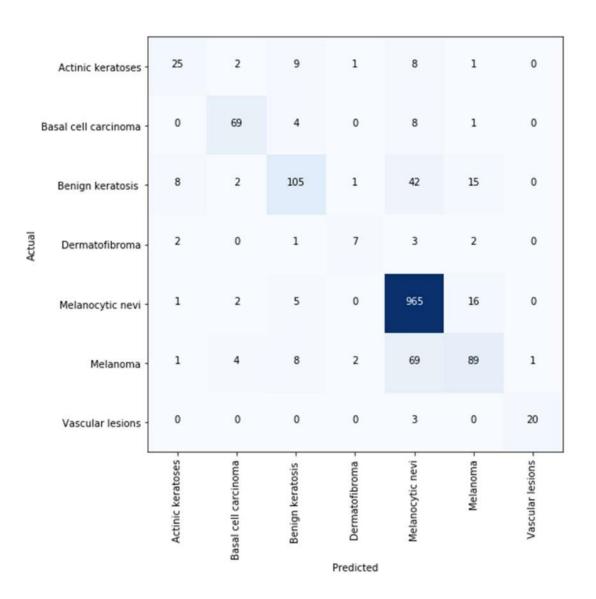




<matplotlib.axes.\_subplots.AxesSubplot at 0x7ff27a102550>



Out[8]:

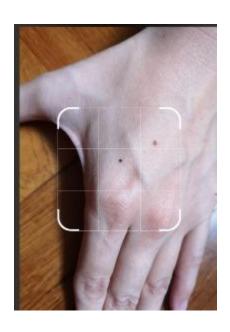


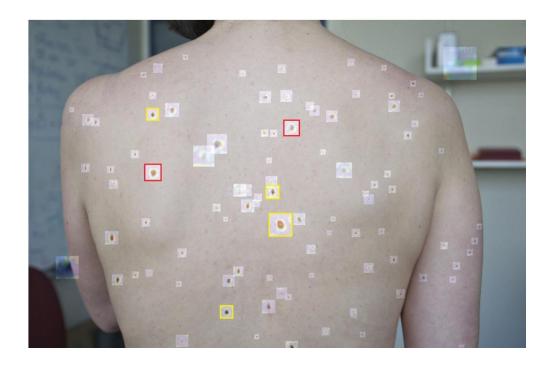
# **RESULTS**

# **9.1 Performance Metrics**

Performance metrics for classification with dement images.

Method	AUC	Specificity	Sensitivity	F1-score
Without segmentation	0.8207	0.9642	0.4748	0.4092
Contextual segmentation	0.8104	0.9652	0.4185	0.3876
Refined contextual segmentation	0.8802	0.9513	0.6141	0.6079





### 10. ADVANTAGES & DISADVANTAGES

# 10.1 Advantages

- → An efficient and quick way to identify the skin problem in absense of a doctor.
- → User friendly interface.
- → User is able to seek possible cure after identification.
- → Portable to use anywhere around the world.
- → Secure data storage in cloud network.

# 10.2 Disadvantages

- → Requires internet connection.
- → A quality camera is required for capturing.

### 11. CONCLUSION

We have shown that even without a large dataset and high-quality images, it is possible to achieve sufficient accuracy rates. In addition, we have shown that current state-of-the-art CNN models can outperform models created by previous research, through proper data pre-processing, self-supervised learning, transfer learning, and special CNN architecture techniques. Furthermore, with accurate segmentation, we gain knowledge of the location of the disease, which is useful in the pre-processing of data used in classification, as it allows the CNN model to focus on the area of interest. Lastly, unlike previous studies, our method provides a solution to classify multiple diseases within a single image. With higher quality and a larger quantity of data, it will be viable to use state-of-the-art models to enable the use of CAD in the field of dermatology.

### 12. FUTURE SCOPE

This implementation of the Structural Co-Occurrence matrices for feature extraction in the skin diseases classification and the pre-processing techniques are handled by using the Median filter, this filter helps toremove the salt and pepper noise in the image processing; thus, it enhances the quality of the images, and normally, the skin diseases are considered as the risk factor in all over the world. Our proposed approach provides 97% of the classification of the accuracy results while another existing model such as FFT + SCM gives 80%, SVM + SCM gives 83%, KNN + SCM gives 85%, and SCM + CNN gives 82%. Future work is dependent on the increased support vector machine's accuracy in classifying skin illnesses, and SCM is used to manage the feature extraction technique

#### 13.APPENDIX

## **GITHUB**

https://github.com/IBM-EPBL/IBM-Project-11717-1659341231

