

# **SKIN DISEASES DETECTION**

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# AGENDA

- 01 Introduction & Problem
- 02 Methodologies
- 03 Related work
- 04 Dataset
- 05 Architecture & Challenges
- 06 Web Application
- 07 Future plans & Conclusion
- 08 References

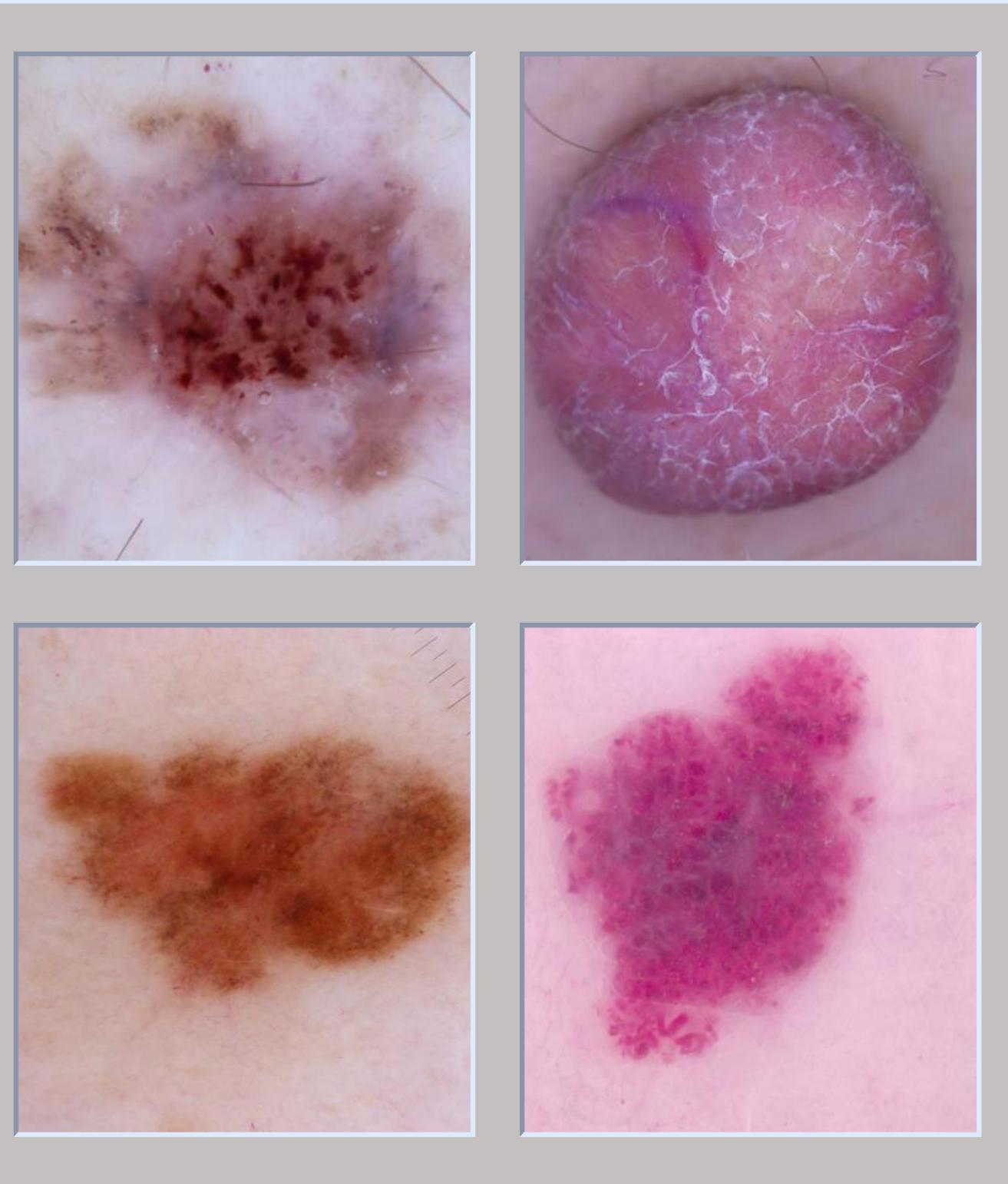
# Introduction

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# Problem

Actinic keratoses



Melanocytic nevus

Dermatofibroma

Vascular lesion

# **Objectives of Project:**



**Explorating  
Appropriate  
Dataset**

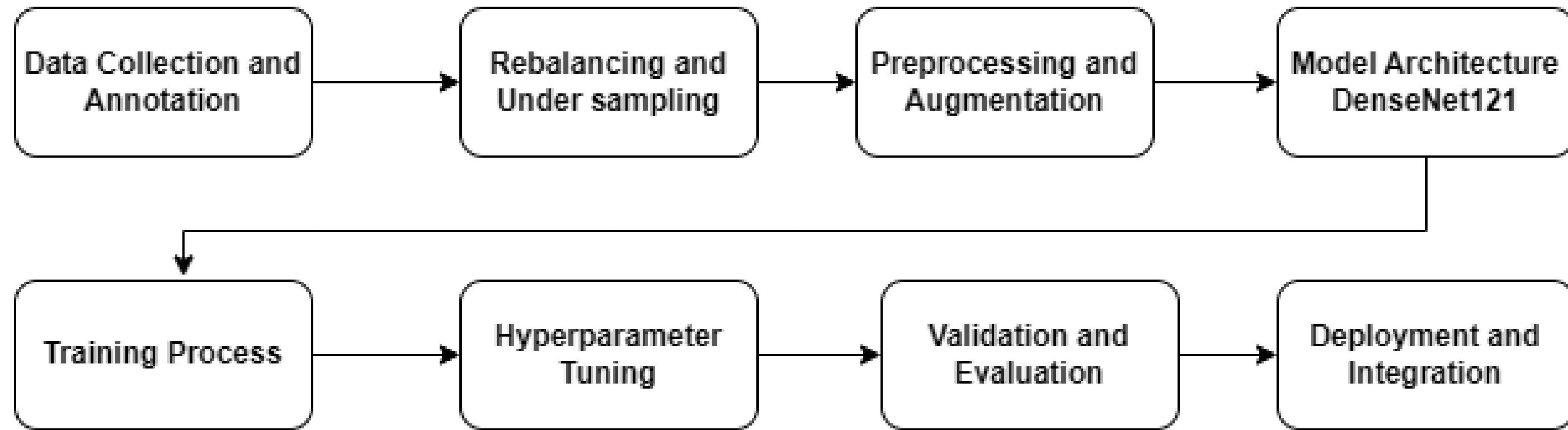
**Building DL  
Model**

**Test and  
Enhance The  
Model**

**Building Web  
App**

**Hosting**

# How



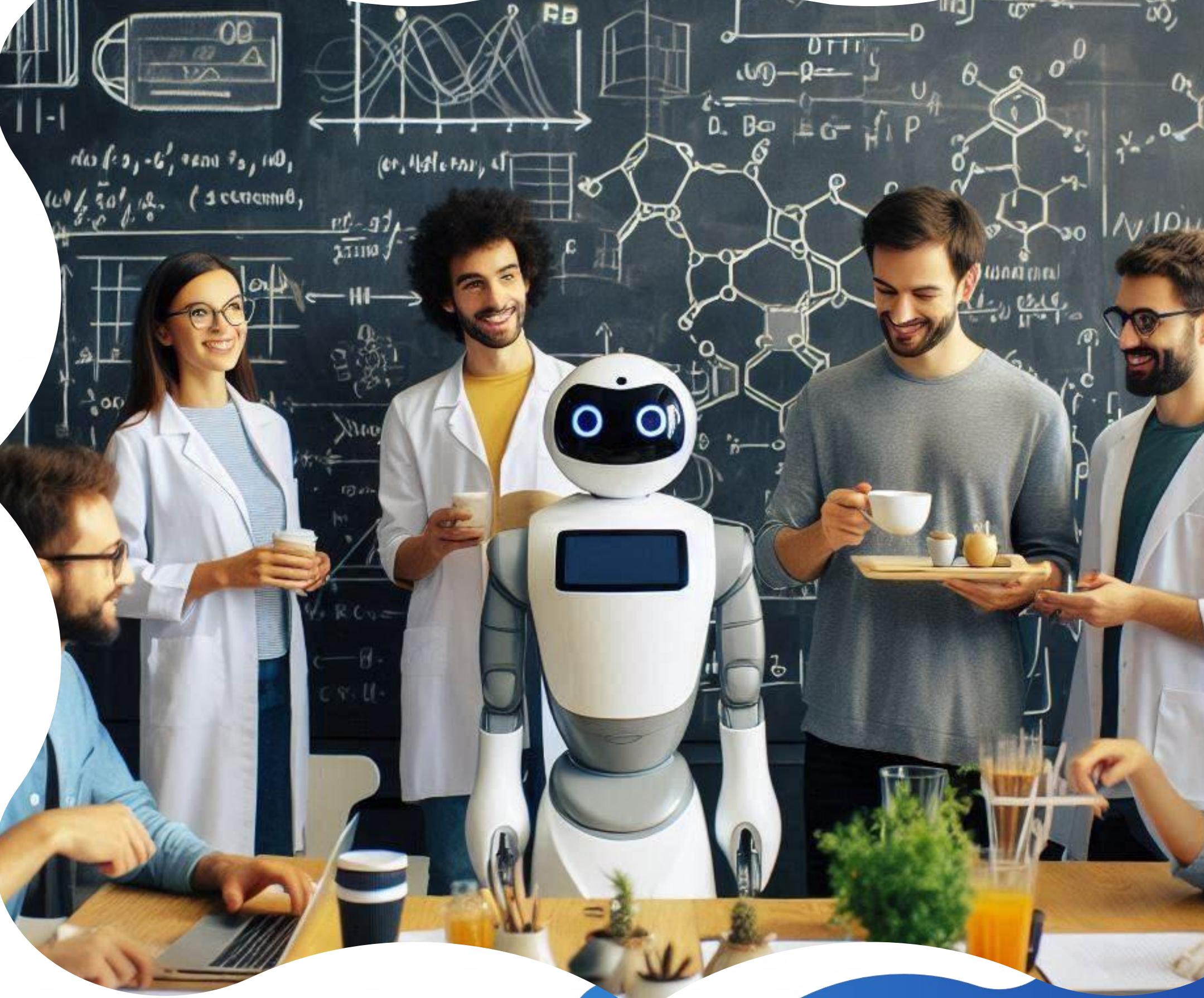
# Methodology

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- We applied insights to address project-specific challenges and optimize our decision-making processes.
- A systematic survey was conducted, analyzing datasets in a structured order, highlighting strengths and weaknesses.
- Making exploration on the algorithms that helped us in emphasizing a methodical evaluation criteria.
- Arranging the results based on priorities in terms of highest accuracy to facilitate decision-making

# Related Work



## References of Skin Diseases with Deep Learning :

Paper	Year	Dataset	Accuracy
<b>Classification of Skin Cancer with Deep Transfer Learning Method</b>	2022	International Skin Imaging Collaboration (ISIC)	97.89%
<b>Multi-Features Extraction Based on Deep Learning for Skin Lesion Classification</b>	2022	ISIC 2019	(92.34% and 91.71%)
<b>Early Detection of Skin Cancer Using Deep Learning Architectures</b>	2019	ISIC-archive	88.20%

# Datasets

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Disease Name	ISIC 2019	HAM10000	PAD-UFES-20	ISIC 2017	Merge
Melanoma	4522	1113	52	521	6208
Melanocytic	12896	6705	244	0	19845
Basal cell	3323	514	845	0	4682
Actinic keratosis	867	327	730	0	1924
Benign keratosis	2624	1099	0	0	3723
Dermatofibroma	239	115	0	0	354
Vascular lesion	253	142	0	0	395
Squamos cell	629	0	192	0	821
Seborrheic keratosis	0	0	235	374	609
Benign nevi	0	0	0	1855	1855
<b>Total</b>	<b>23127</b>	<b>11711</b>	<b>5578</b>		<b>40416</b>
		<b>17289</b>			

# Priority Criteria

ISIC 2019



HAM 10000



PAD-UFES-20



ISIC 2017



**Availability**



**Aim to facilitate training neuron network**



**Suitable number of skin diseases.**



**Large number of images in each disease.**

# Unused Datasets

## DermNet

1. Num\_images is very small
2. Aim to archive to medical.
3. lack of Organization.

## Derm7pt

1. Num\_images is very small.
2. Diseases don't intersect with the majority of available diseases.

## BCN20000

1. Unavailable independently.
2. Embedded with ISIC 2019.

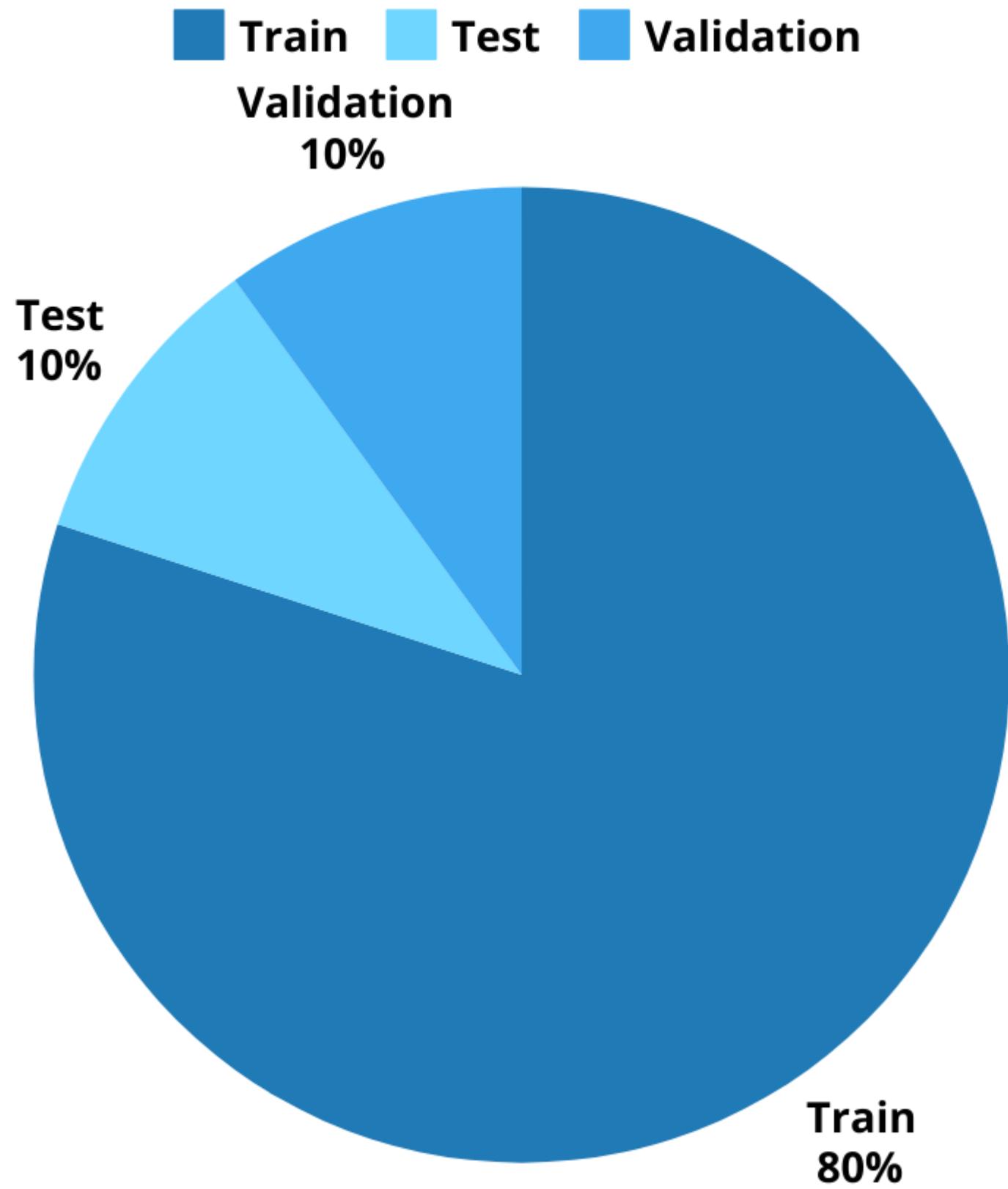
## PH2

1. Num\_images is very small.
2. Diseases are very general.

## Dermofit

Unavailable

# DataSet from :

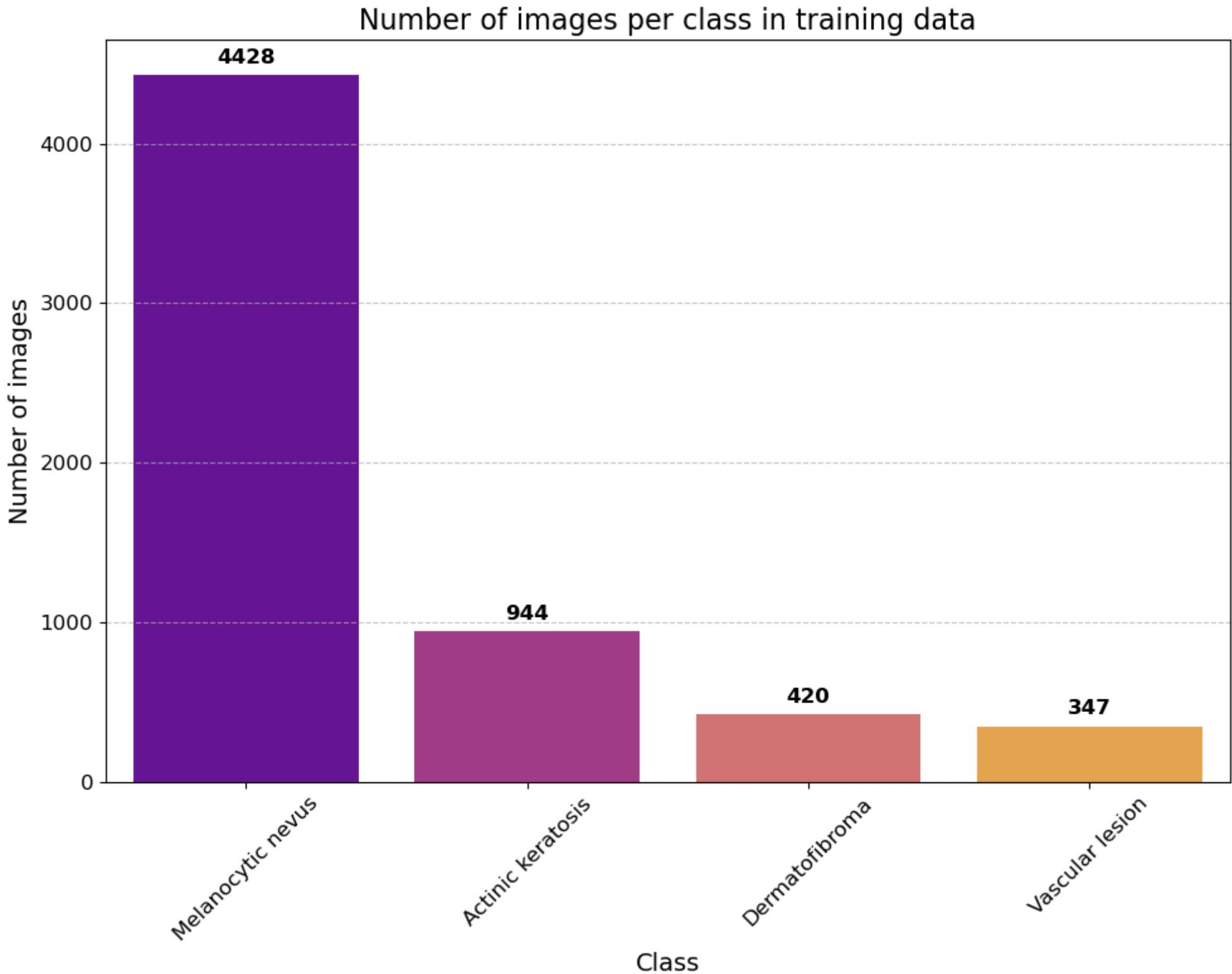


# Preprocessing



# Under sampling :

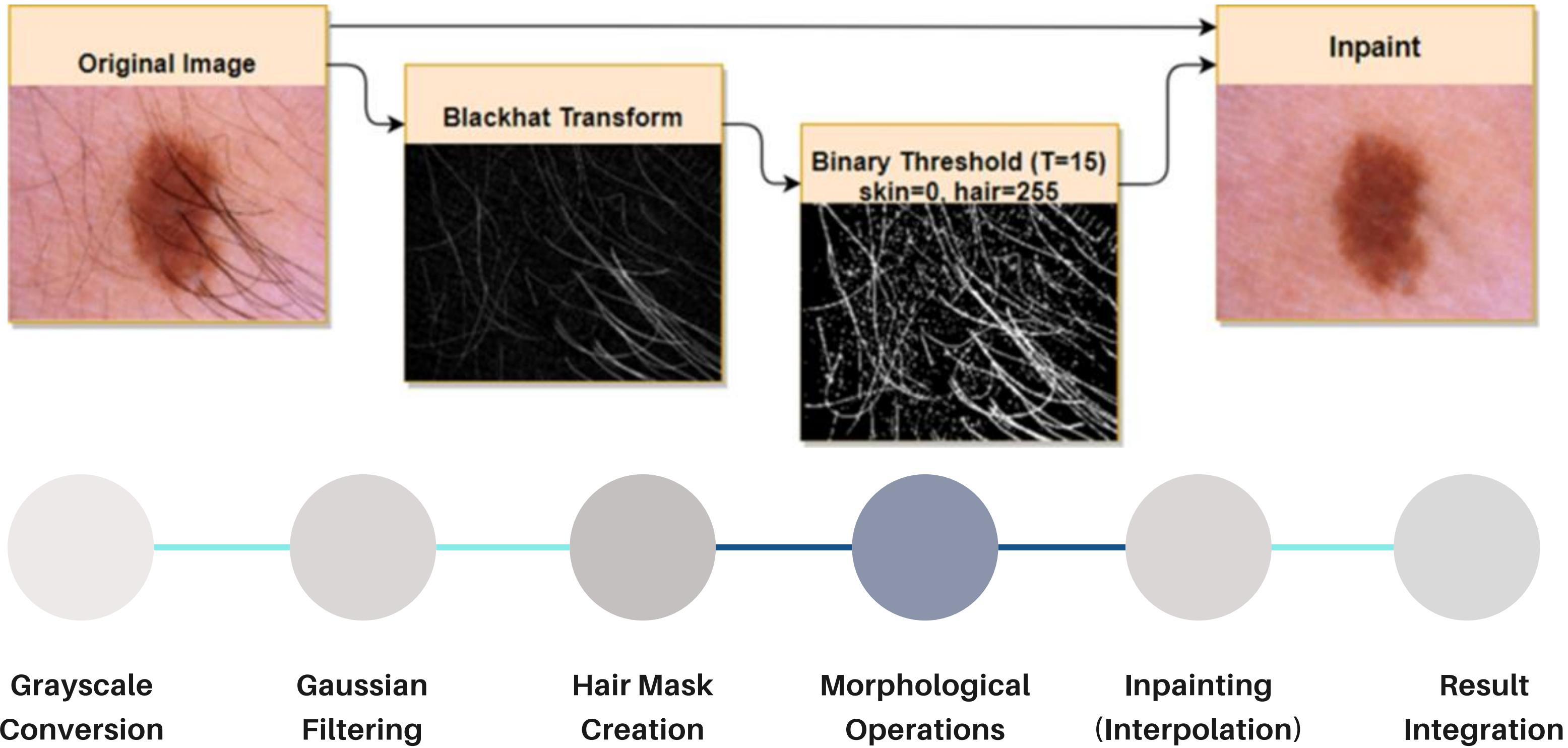
- An exaggerated increase for Melanocytic
- There are few remaining diseases
- Reduce Melanocytic From 11k to 4k



# Data Augmentation

<i>rescale</i>	1.0/255
<i>rotation range</i>	40
<i>width shift range</i>	0.2
<i>height shift range</i>	0.2
<i>shear range</i>	0.2
<i>zoom range</i>	0.2
<i>fill mode</i>	nearest
<i>horizontal flip</i>	true
<i>feature wise std normalization</i>	true

# Dull razor Algorithm

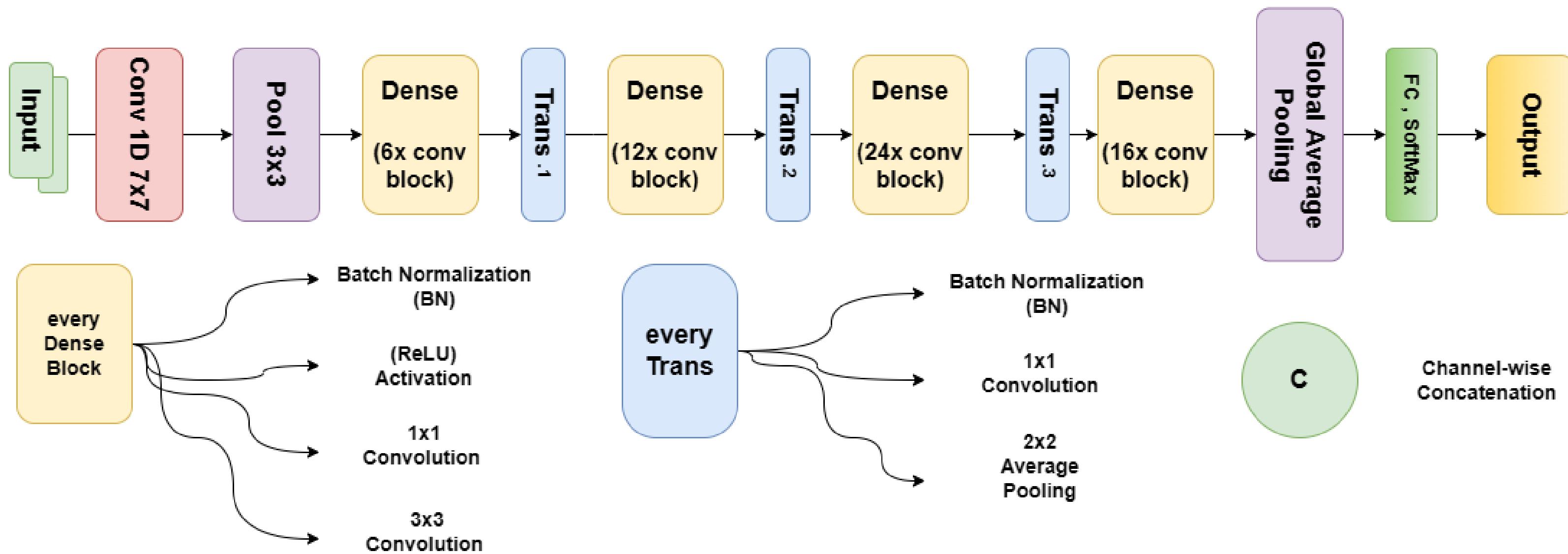


# Proposed Final Model

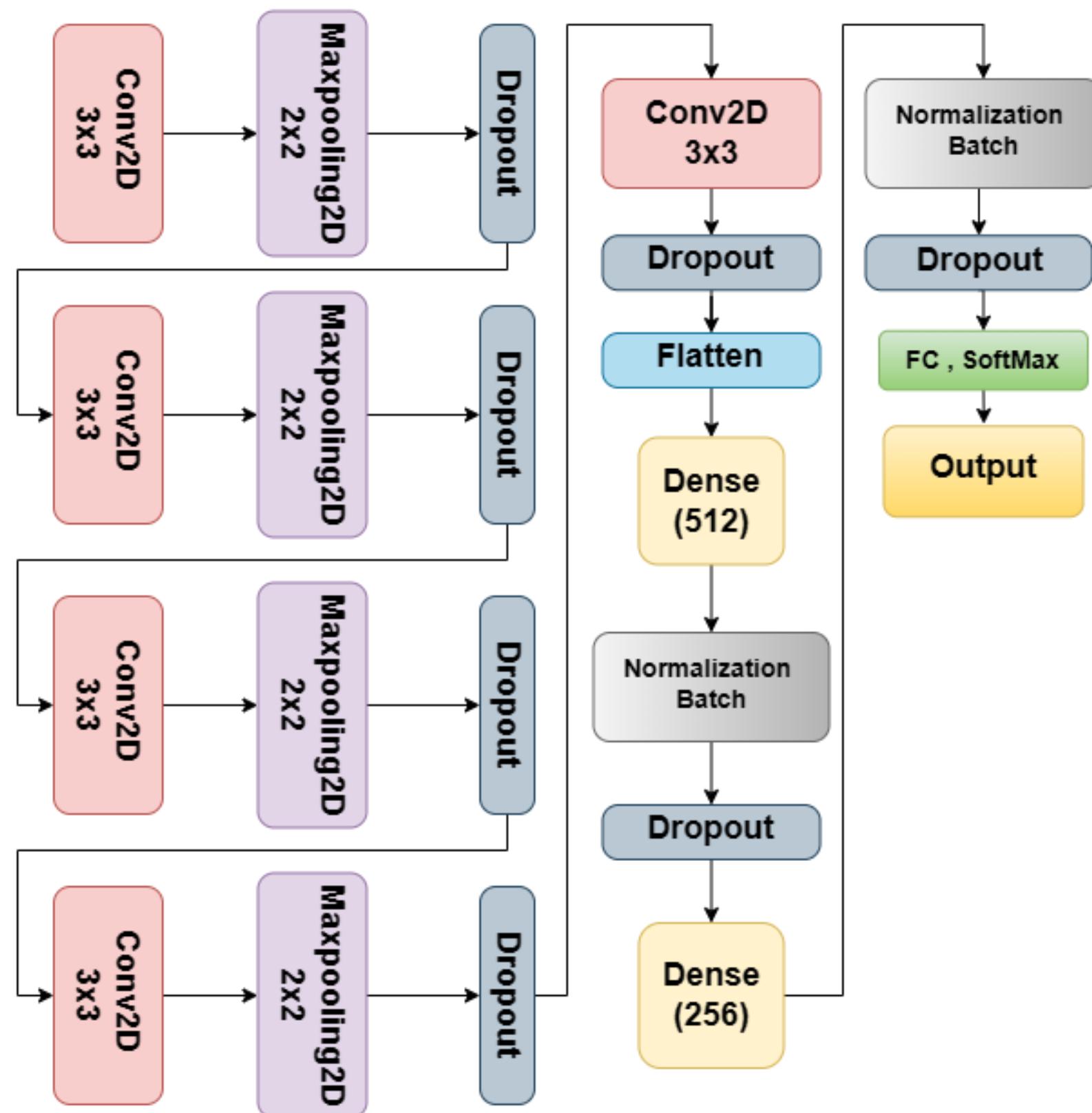


# Architecture of DenseNet121

The Base Model with pre-trained Dataset ImageNet



# Custom Layers:



- Conv2D and Dense use activation Function Relu
- MaxPooling2D and Conv2D padding is same
- Dropout (0.5/0.3)

# Freezing Layers:

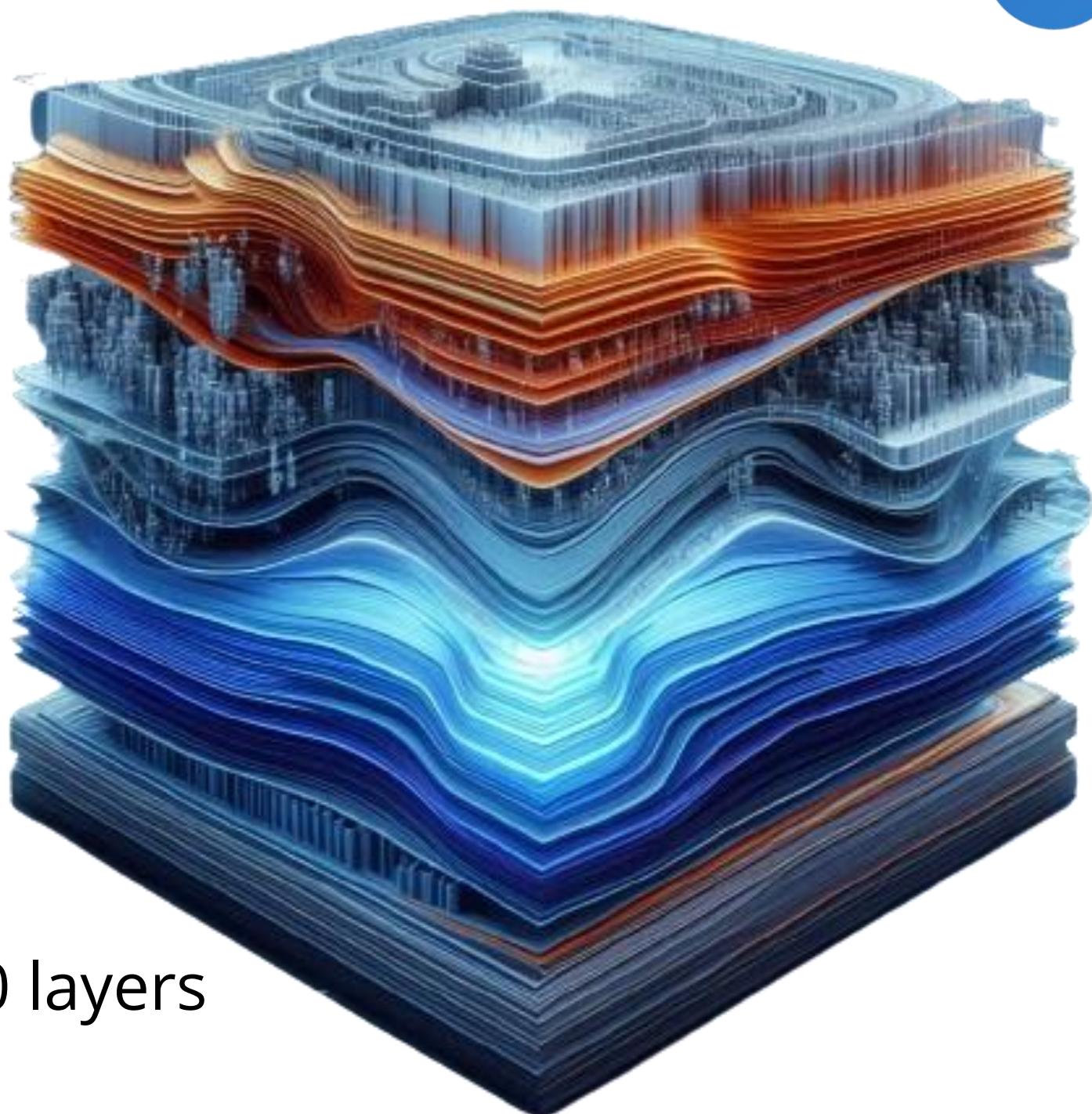
the combination of freezing and selectively unfreezing layers is a key technique in transfer learning.

**Preserve Learned Features**

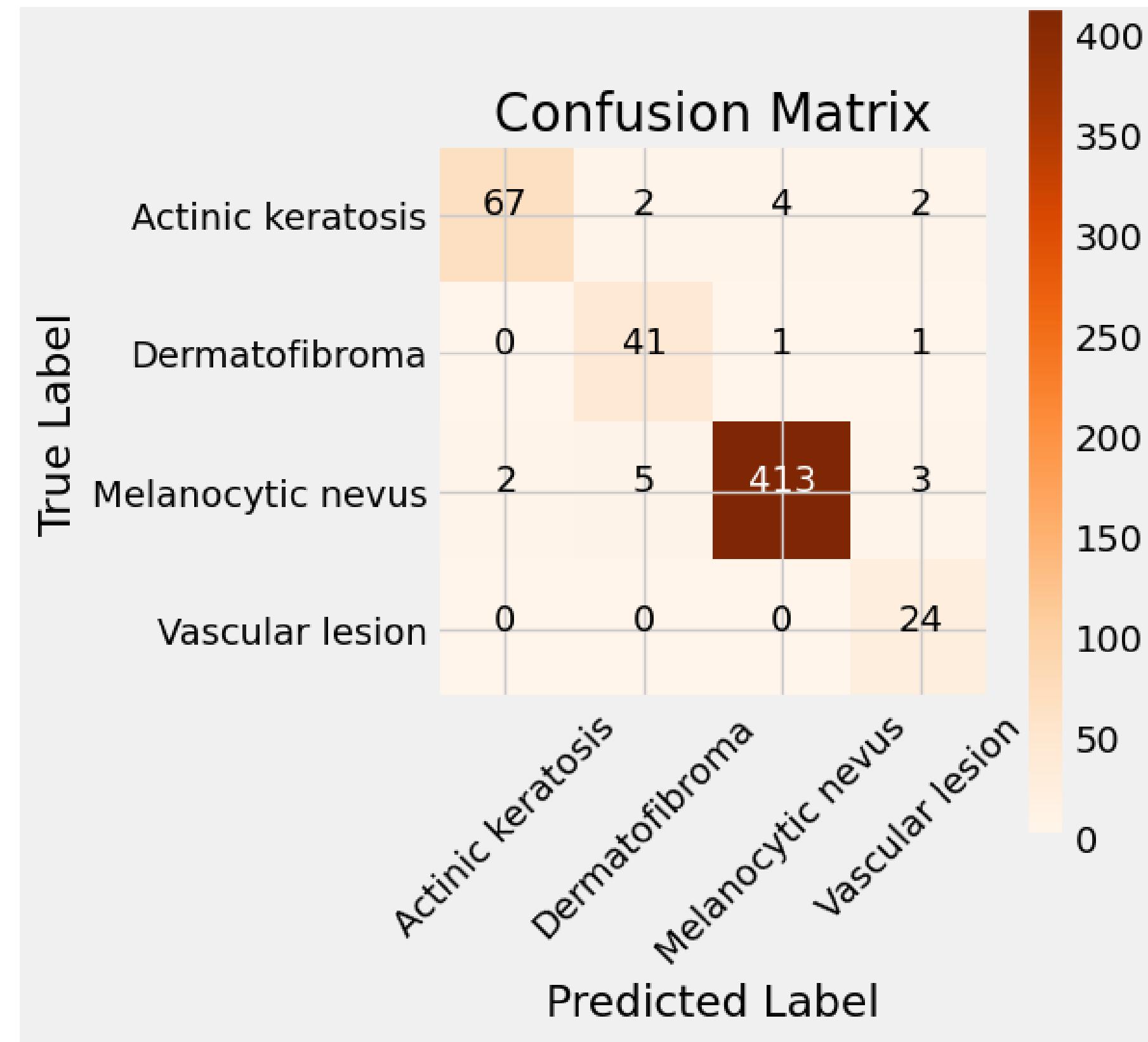
**Adapt to New Task**

**Efficient Training**

Unfreeze the top 30 layers



# Confusion Matrix:

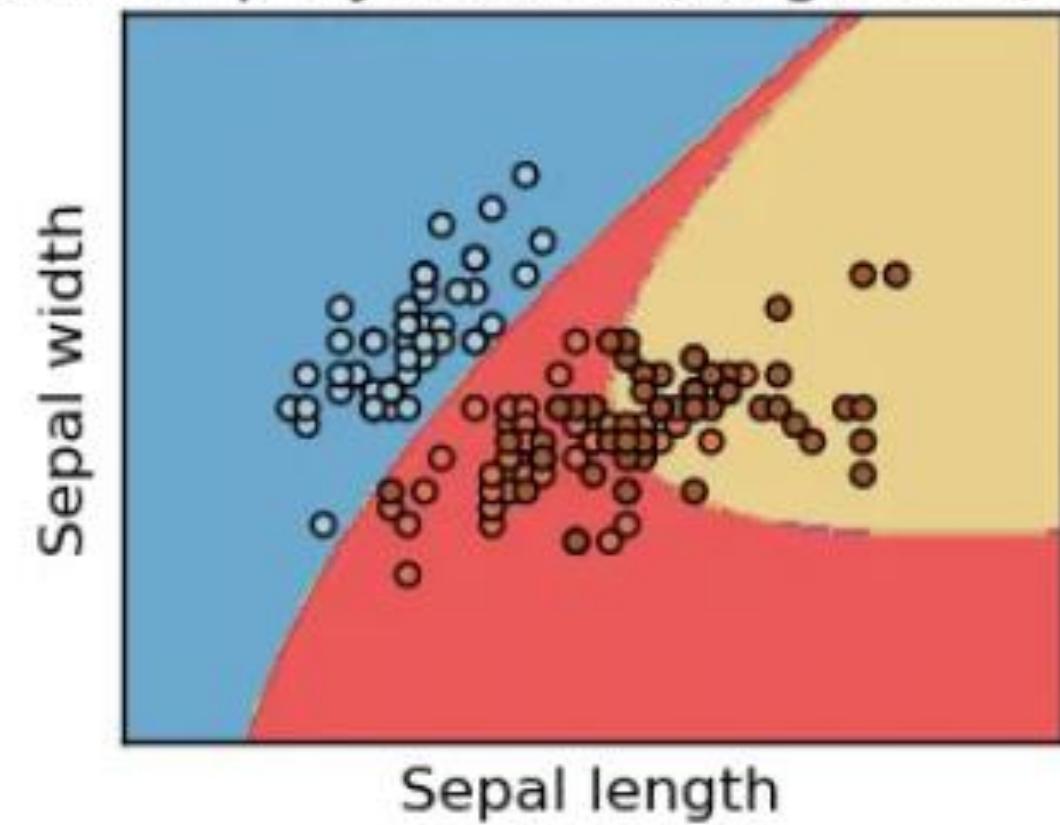


# Classification Report:

	Precision	Recall	F1-Score	Support
Actinic keratosis	0.97	0.89	0.93	75
Dermatofibroma	0.85	0.95	0.9	43
Melanocytic nevus	0.99	0.98	0.98	423
Vascular lesion	0.8	1	0.89	24
<b>Accuracy</b>			0.96	565
<b>Macro avg</b>	0.9	0.96	0.93	565
<b>Weighted avg</b>	0.97	0.96	0.97	565

# Svc Method :

SVC with polynomial (degree 3) kernel

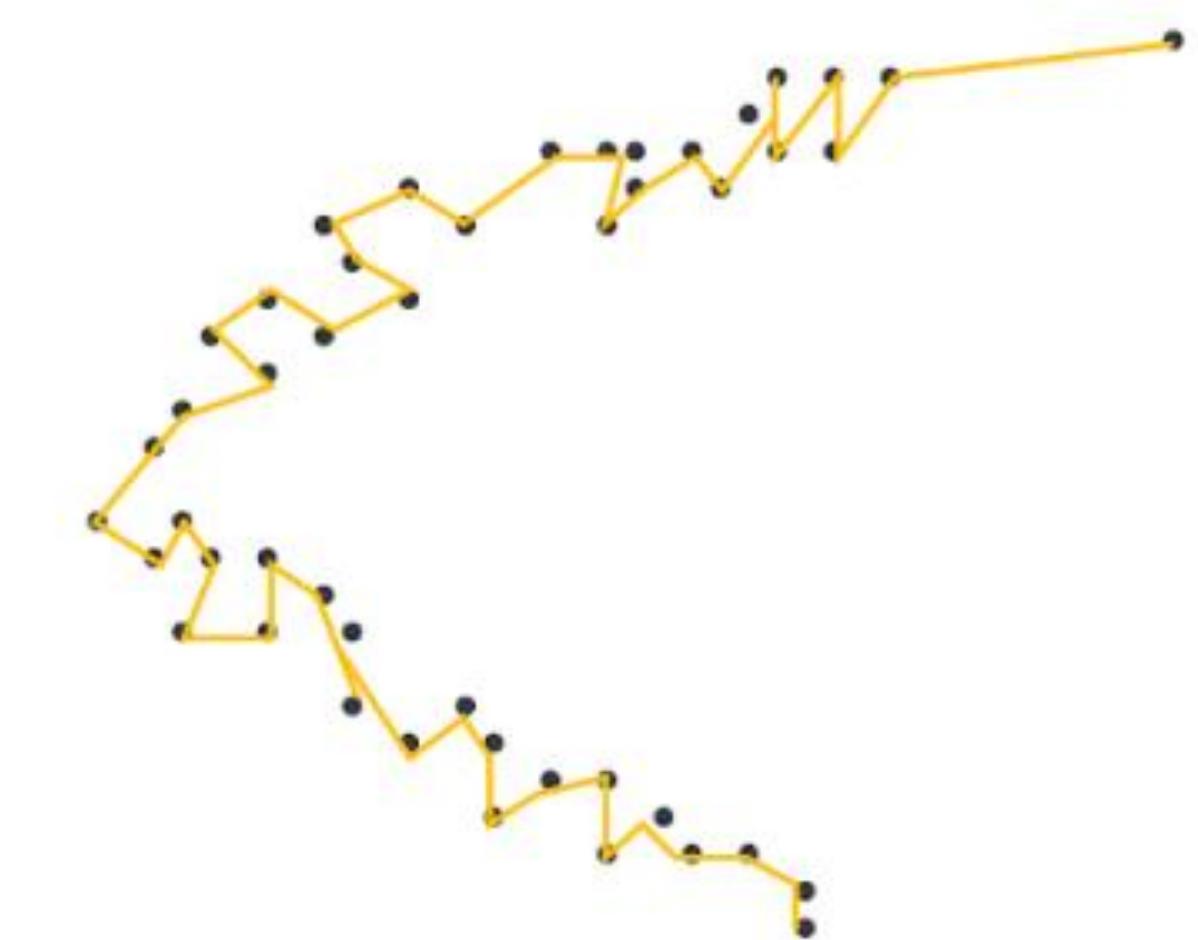


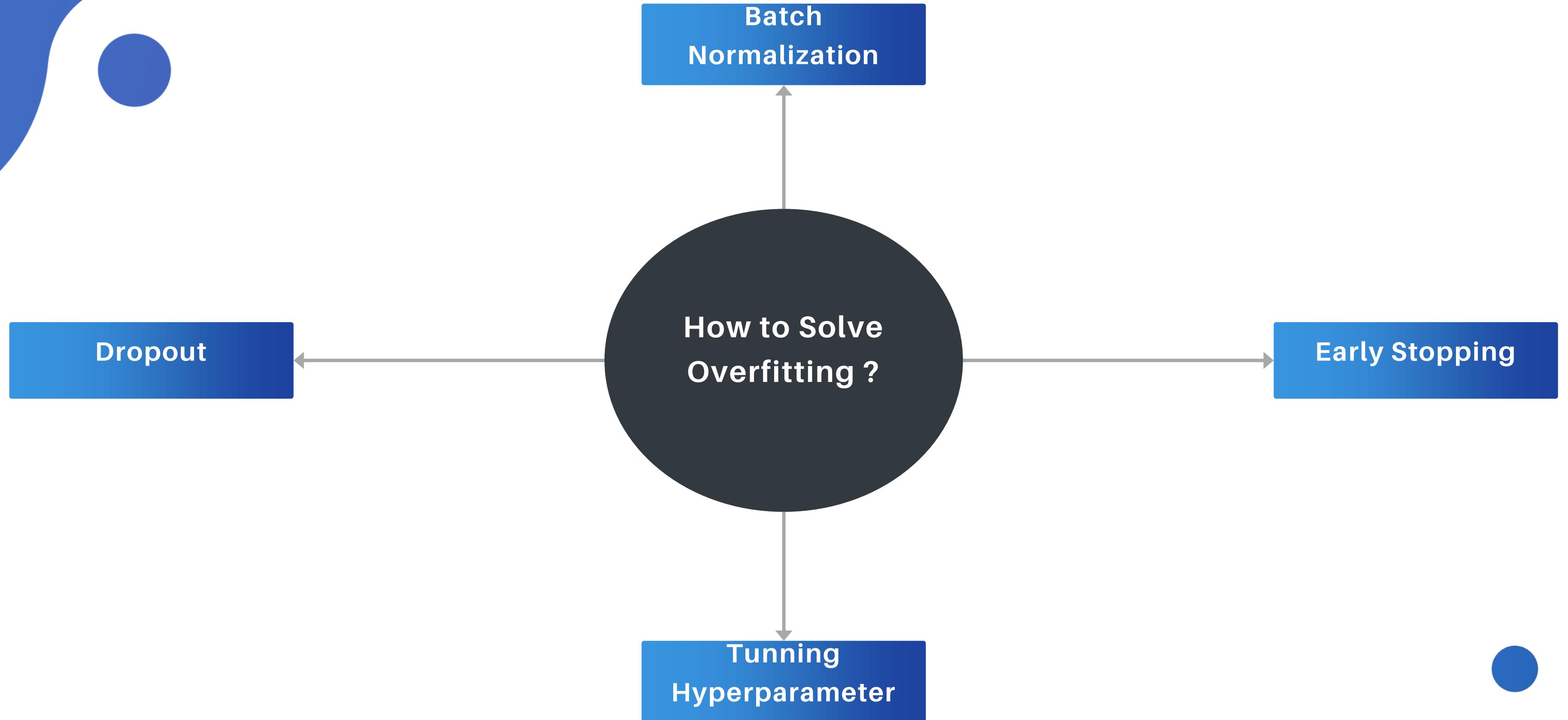
```
1 test_features, test_labels = extract_features(model, test_generator)
2
3 svc_model = Pipeline([
4     ('scaler', StandardScaler()),
5     ('svc', SVC(kernel='poly', degree=3))
6 ])
7 svc_model.fit(test_features, test_labels)
```

# What is Overfitting?

Overfitting is a modeling error in machine learning and statistics where a model is too complex and fails to generalize to new data.

It occurs when a model is too tailored to the training data, memorizing it instead of learning true patterns, resulting in poor performance on test data or real-world applications.





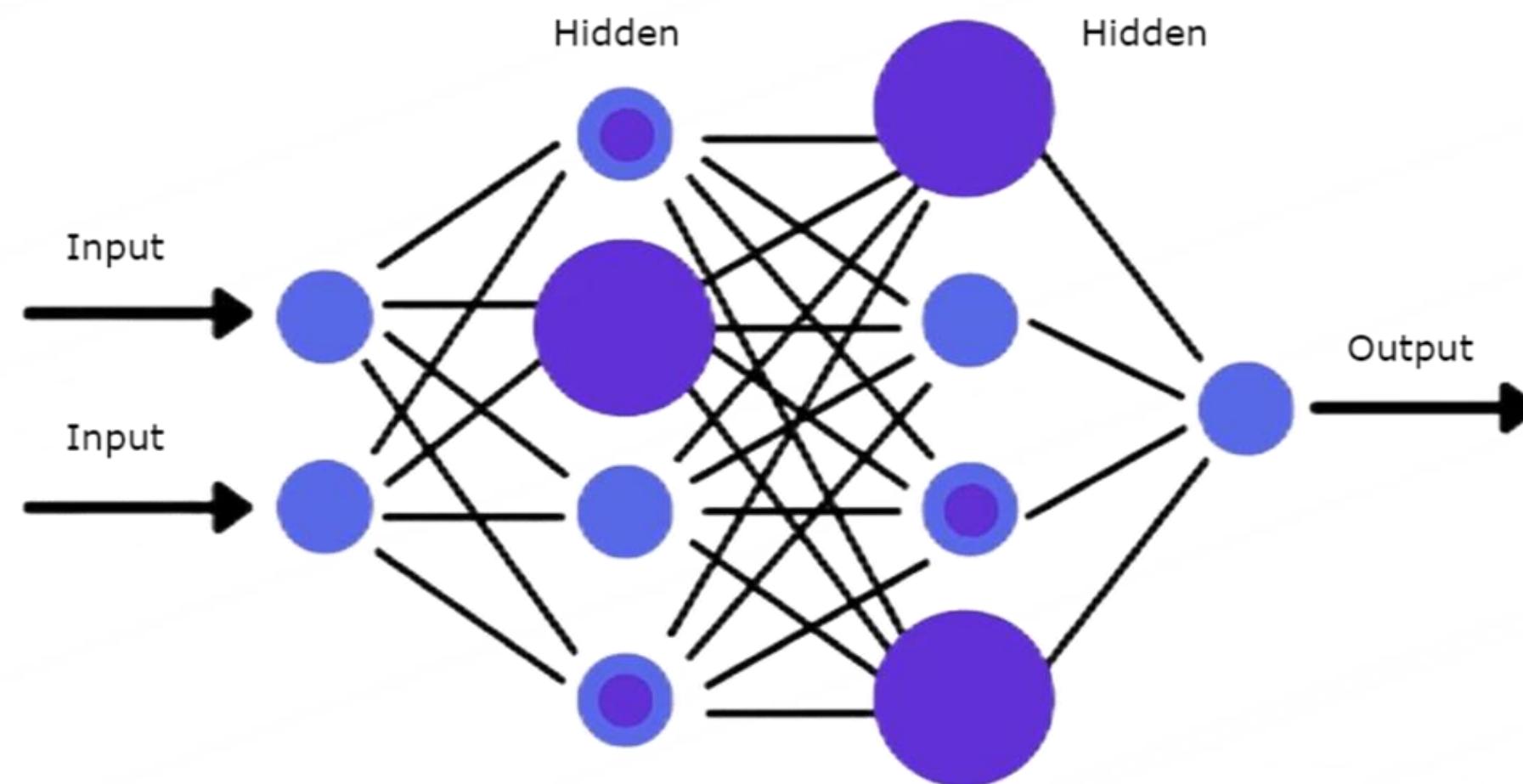
# Hyperparameter

:

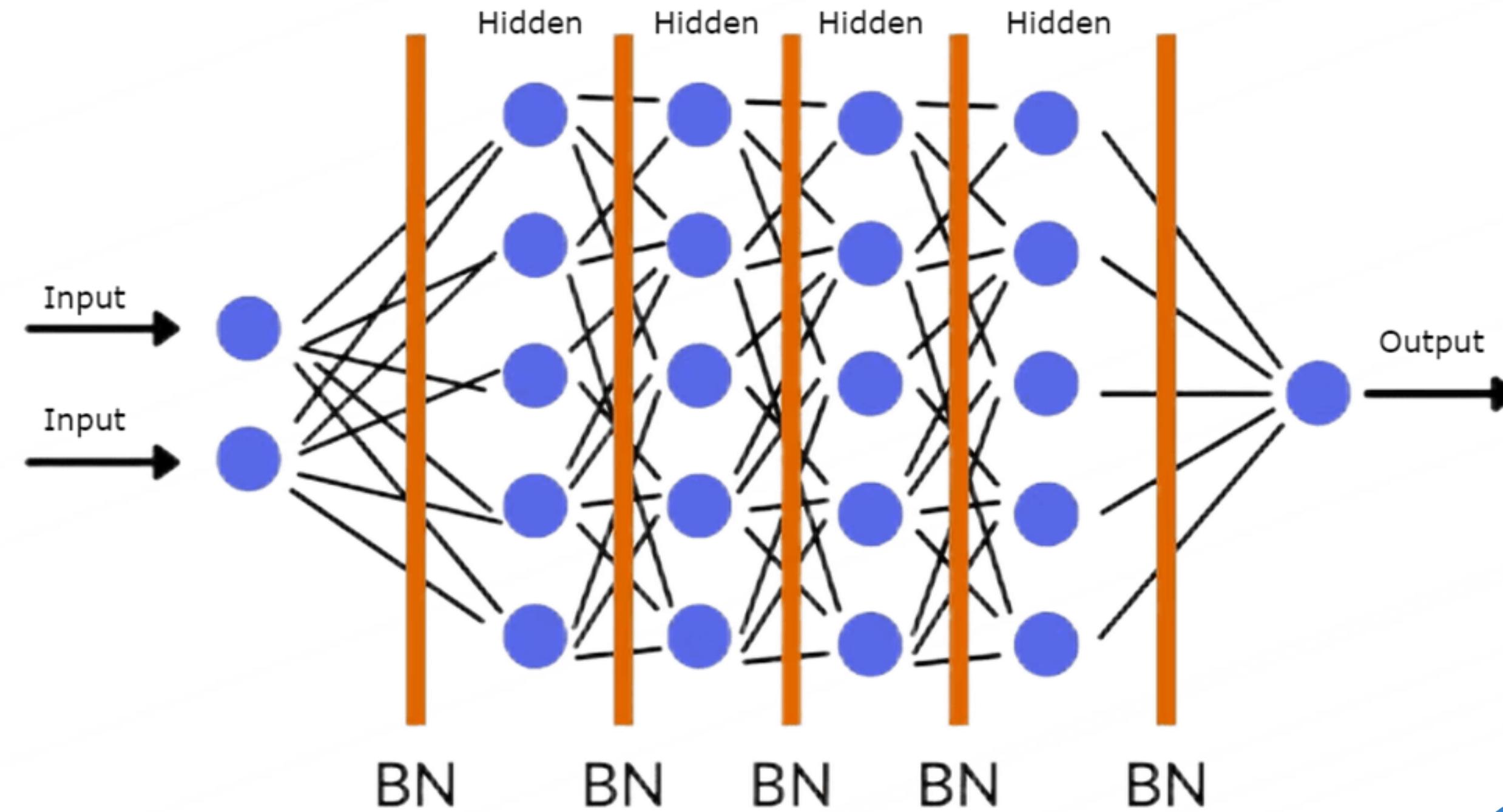
<b>loss function</b>	categorical_crossentropy
<b>activation function</b>	Relu
<b>dropout rates</b>	0.5   0.3
<b>epochs</b>	100
<b>batch size</b>	100
<b>image size</b>	224 x 224
<b>learning rate</b>	0.001

# Batch Normalization

Batch normalization is a deep neural network training technique that reduces internal covariate shift by normalizing inputs to a mean of zero and variance of one.



# During Batch Normalization



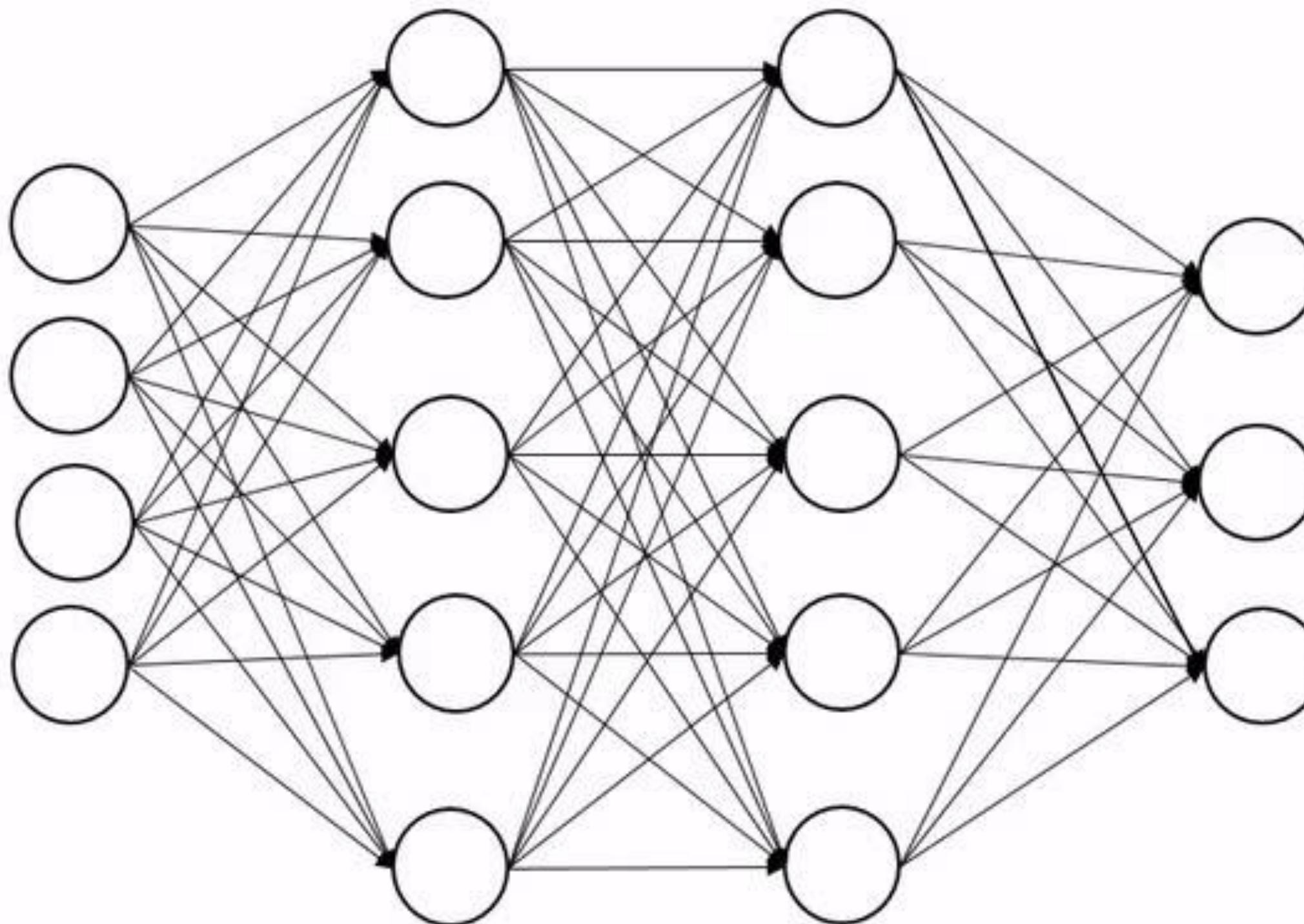
# Dropout

## **Randomly Disable Neurons:**

By randomly ignoring neurons during training

During training, the remaining neurons are scaled up to maintain expected output values, such as multiplying the output of the remaining neurons by 2 if half are dropped out.

# Dropout



# Early Stopping

Method to avoid [overfitting](#).

Early stopping involves terminating training shortly once the validation error is at its lowest level.

Many variables are used in early stops :

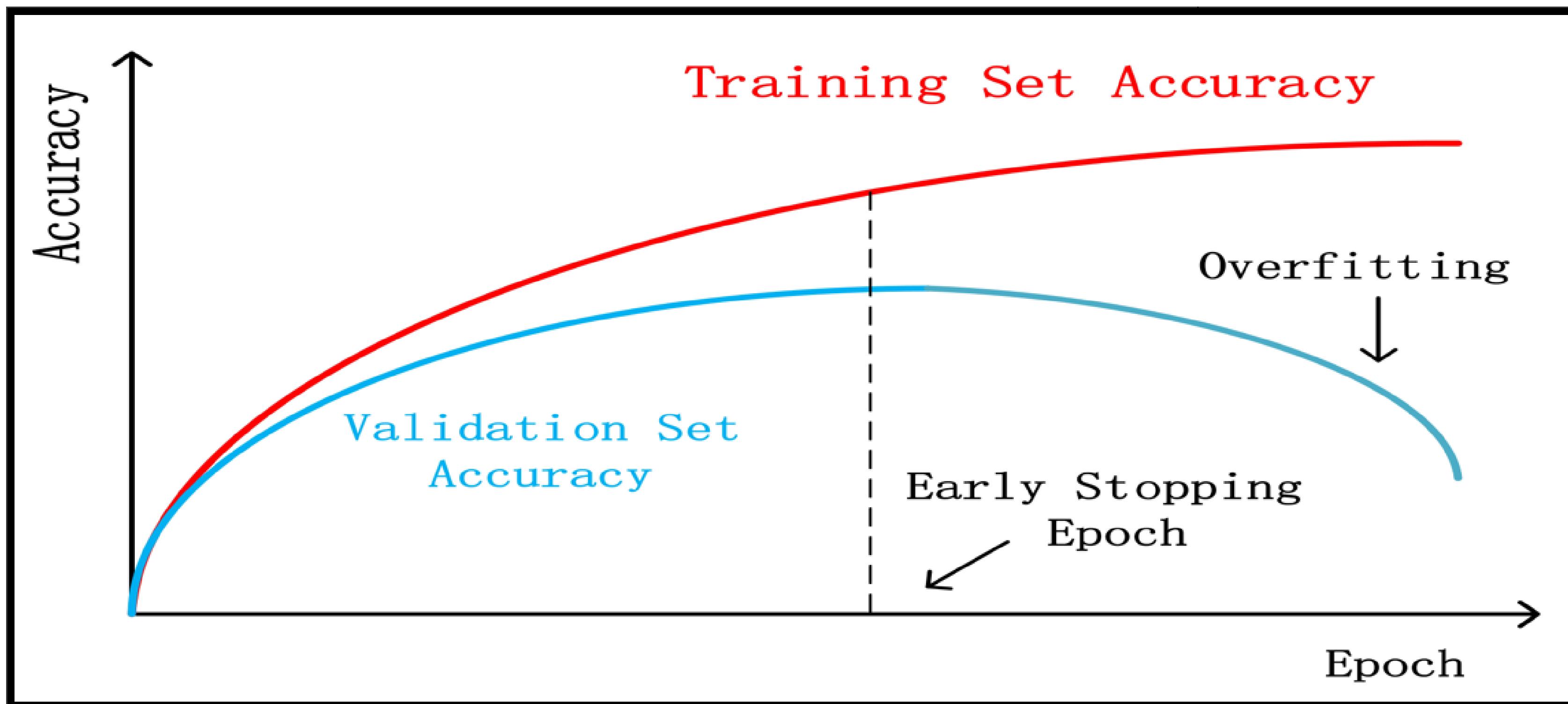
***monitor= 'loss'***

***mode= 'min'***

***patience= 10***

***min\_delta= 0.001***

# Early Stopping



# Experiments

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# Results of Experiments: 2 April

<b>epochs</b>	10
<b>Input size</b>	224,224
<b>Reduce lr</b>	no
<b>Dropout</b>	0.3   0.5
<b>Unfreeze Layers</b>	30 top layer from base
<b>early_stop</b>	no
<b>Test acc</b>	0.860
<b>Test Loss</b>	0.717

# Results of Experiments: **3 May**

<b>epochs</b>	15
<b>Input size</b>	250,250
<b>Reduce lr</b>	yes (factor=0.1,patience = 15)
<b>Dropout</b>	0.5   0.5
<b>Unfreeze Layers</b>	top 30 layer
<b>early_stop</b>	no

<b>Test acc</b>	0.914
<b>Test Loss</b>	0.422

# Results of Experiments: 15 April after using SVC method

<b>epochs</b>	100
<b>Input size</b>	224,224
<b>Reduce lr</b>	no
<b>Dropout</b>	0.5   0.3
<b>Unfreeze Layers</b>	30 from top
<b>early_stop</b>	(monitor='loss', mode='min',patience=10)

<b>Test acc</b>	0.959	<b>Cubic SVM Test acc</b>	0.9
<b>Test Loss</b>	0.137	<b>Cubic SVM Test Loss</b>	0.09999

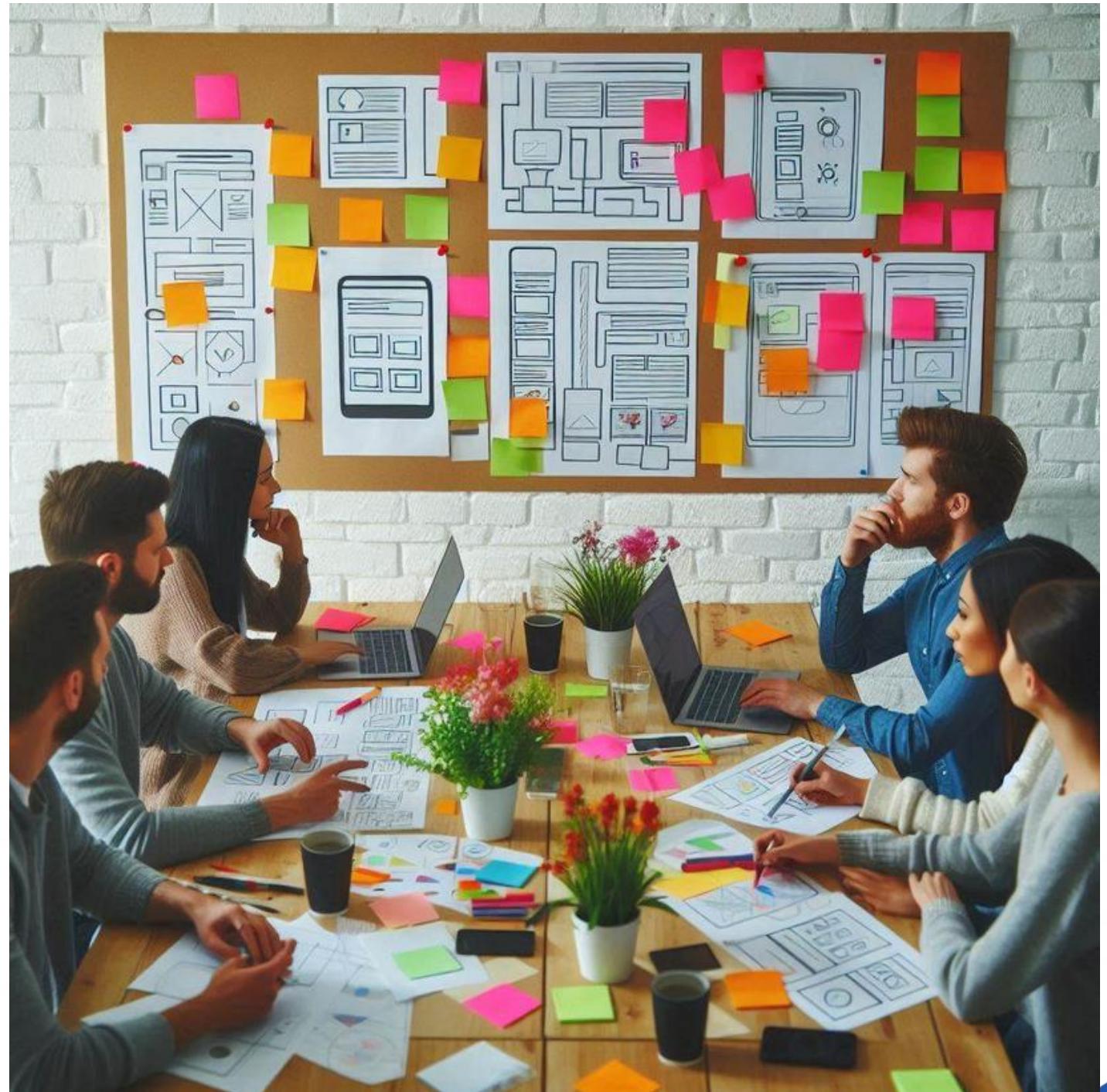
# Results of Experiments: 30 April after reduce Melanocytic to 4k

<b>epochs</b>	100
<b>Input size</b>	224,224
<b>Reduce lr</b>	no
<b>Dropout</b>	0.5   0.3
<b>Unfreeze Layers</b>	30 from top
<b>early_stop</b>	(monitor='loss', mode='min',patience=10)

<b>Test acc</b>	0.964
<b>Test Loss</b>	0.127

<b>Cubic SVM Test acc</b>	0.99
<b>Cubic SVM Test Loss</b>	0.010000

# Web Application



# UI/UX

Skin  
Disease

Header #1



Try Now

Home App About us

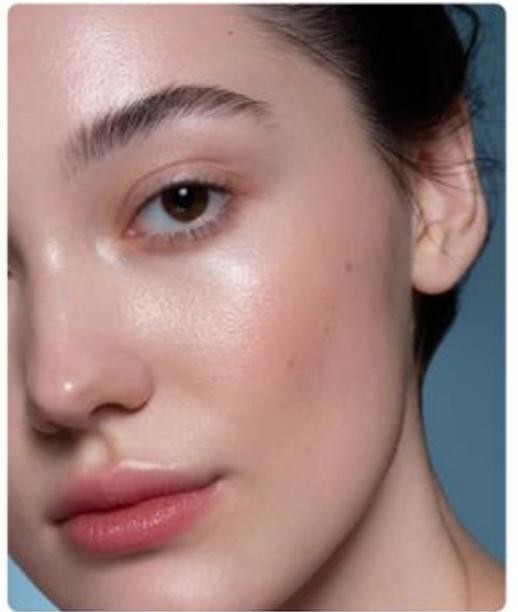


Skin  
Diseases

Caring for you &  
your family

Our target help you determine  
the stage of your disease.

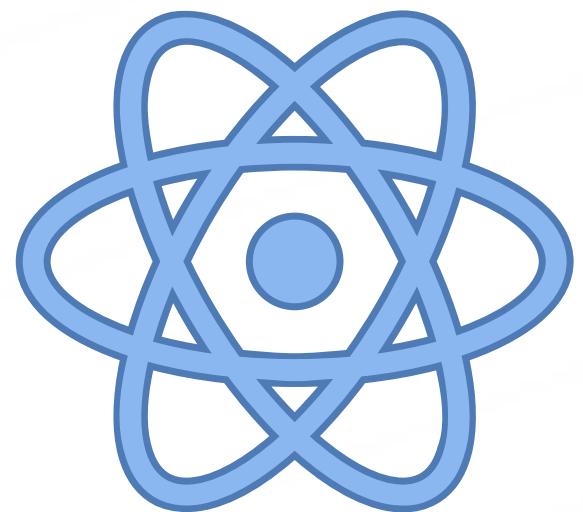
TRY NOW



Full design in Figma

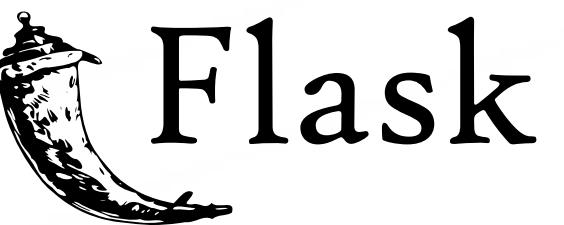
# Application Stack

Front-end



React

Back-end



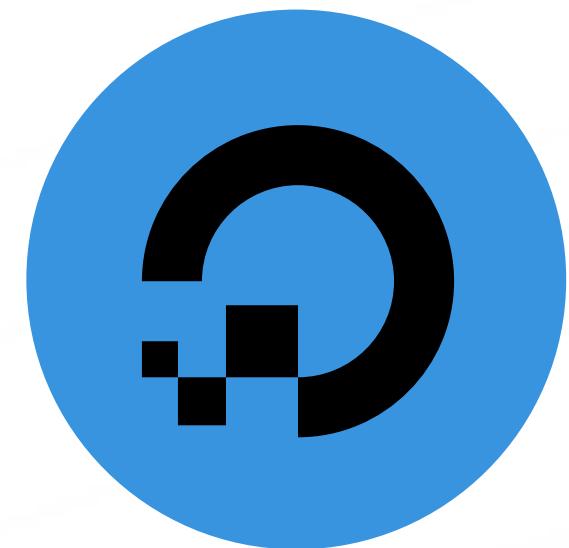
Flask

Webserver

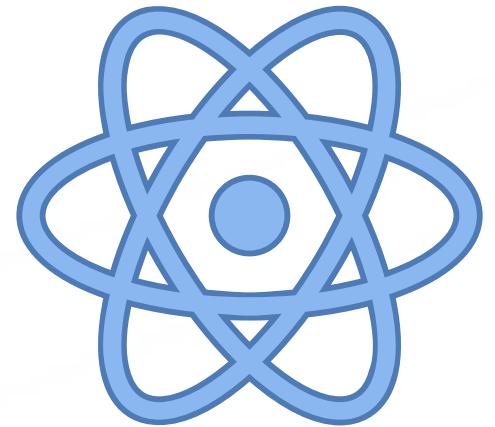


Nginx

Cloud Platform



Digital Ocean



## Data flow

- API Requests
  - Fetch data from backend API
  - Handle responses and errors
- State Management
  - Update components based on state changes

## Libraries

- axios
- react-dropzone
- react-slideshow

## Tools

- eslint for code linting
- Vite for module management



API Endpoint

AI implementation

Server response

```
@app.route('/upload', methods=['POST'])
def predict():
    if 'file' not in request.files:
        return jsonify({'error': 'No file'}), 400

    file = request.files['file']
    img_data = file.read()

    img = image.load_img(io.BytesIO(img_data), target_size=(224, 224))
    img = image.img_to_array(img)
    img = np.expand_dims(img, axis=0)
    img /= 255.0
    prediction = model.predict(img)
    predicted_class = np.argmax(prediction, axis=-1)
    label = class_labels[predicted_class[0]]
    print(label)

    return jsonify({'prediction': label})
```

# Demo

Discover Dermatology

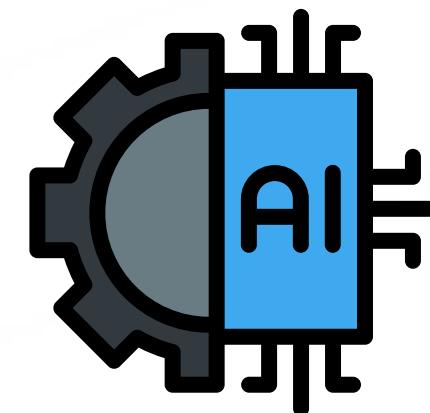


## Melanocytic

Melanocytic lesions are common and can be either benign or malignant. They arise from melanocytes, which are pigment-producing cells.

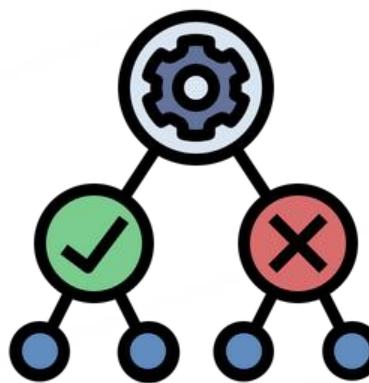


# Computational time



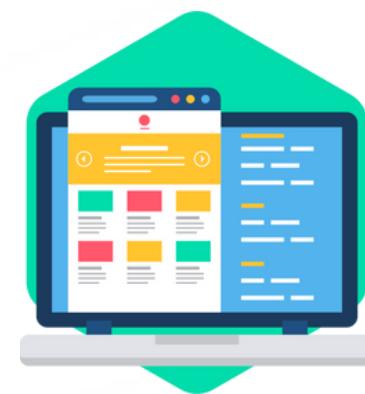
**Training  
model**

4 hour,  
26 minutes,  
and 40  
seconds



**Testing model**

32  
second



**Testing Web  
App**

About 5 seconds

# Future plans & Conclusion

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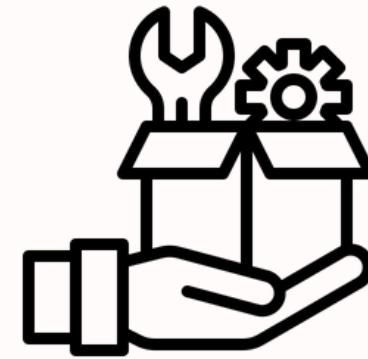
# Conclusion

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- The DenseNet-121 algorithm was chosen due to its proven ability to capture intricate features and efficiently manage complex image data.
- The integration of a Cubic SVM classifier further enhanced the accuracy, achieving a test accuracy of 0.99 in the best experimental setup.
- Effectively resolved the project's several challenges.
- Create a successful Web application that will allow us to sell it with additional work and marketing.

# Future Plans

## What we hope to achieve :



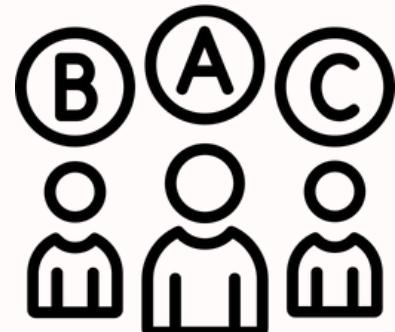
### Expanding service

Increasing the number of dermatologists on the web Application and expanding the service regions .



### Add User Profile

Keep track of their personal data, medical history, and previous diagnoses .



### Categorize Users Groups

Creating 2 categories of users by dividing membership into patients and doctors

# Future Plans

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## What we hope to achieve :



### **contribution**

We aim to publish a comprehensive research paper on our skin disease classification accuracy rate of 0.99, presenting new results in dermatological diagnostics.

# References

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<https://www.kaggle.com/datasets/kmader/skin-cancer-mnist-ham1000>

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<https://arxiv.org/pdf/1512.03385.pdf>

**THANK'S  
FOR  
WATCHING**

