

SIIM-ISIC Melanoma Classification

Introduction:

- Dermatologists analyze lesion images to identify melanoma in patient's mole
- Early detection of melanoma can be cured with appropriate treatment
- Efficient computer vision algorithm can reduce visual error and time spent in the manual classification of lesion images
- Society for Imaging Informatics in Medicine (SIIM) pursuit of developing image analysis tools that automate melanoma diagnosis

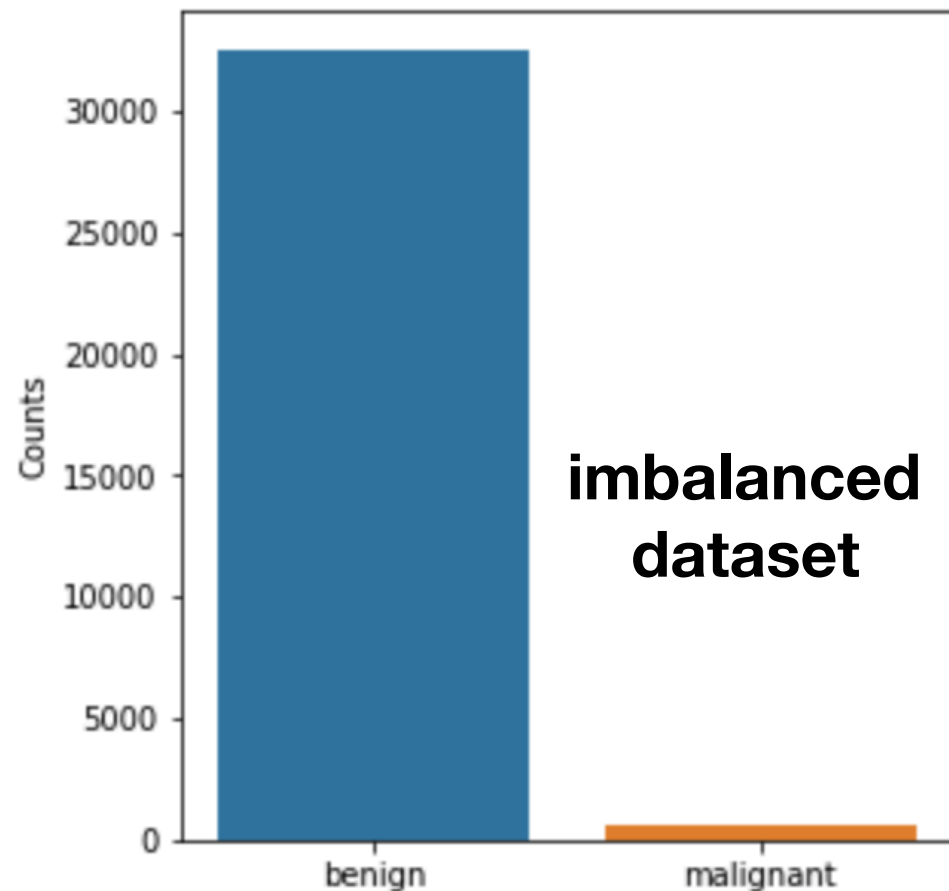
Objective:

- Using ISIC archived dermoscopic images (available in kaggle), develop model to classify lesion images to a benign or malignant class

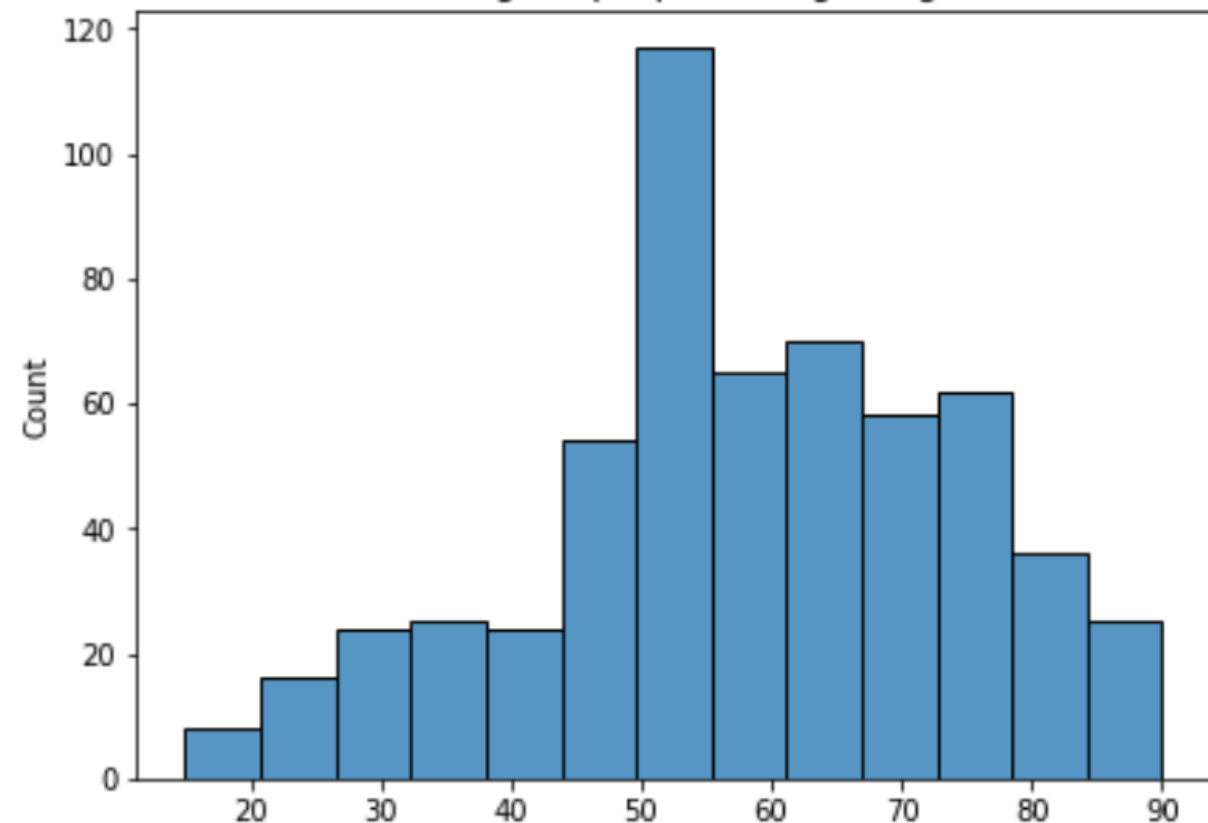
Dataset

- Kaggle's competition SIIM-ISIC Melanoma classification 2020
- 33,216 images with metadata such as patients id, sex, age, anatomic site, diagnosis, lesion label and target columns referring benign or melanoma
- 32,542 benign images with 584 (1.76%) melanoma images
- 2056 patients whose number of images ranges between 2-115
- Malignant observed across all ages of patient

No of images



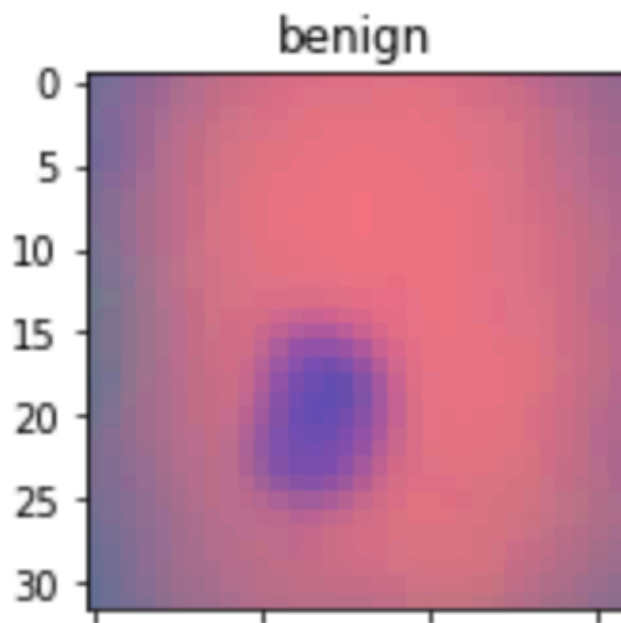
Distribution of age of people having malignant leison



Signs of Melanoma

- Asymmetric mole with irregular and uneven edges
- Large brownish spot (diameter >6 mm)
- Mole that changes in color, size or feel or that bleeds
- Lesion that itches or burn

Benign image



Melanoma images

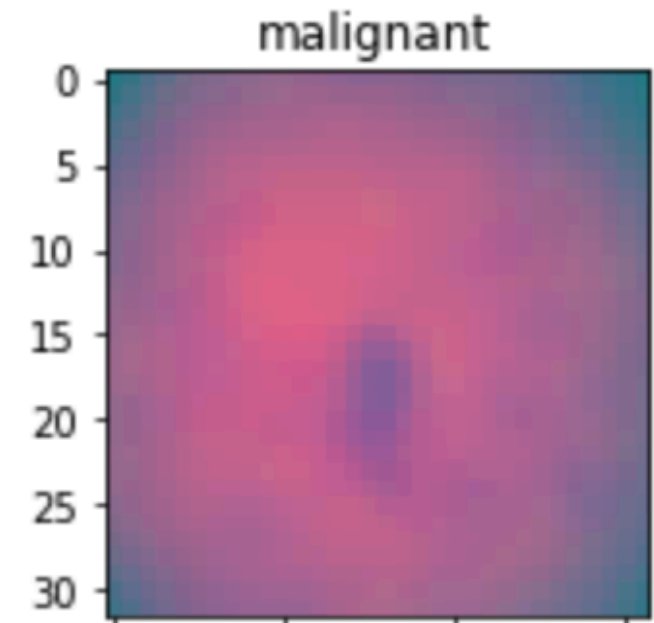
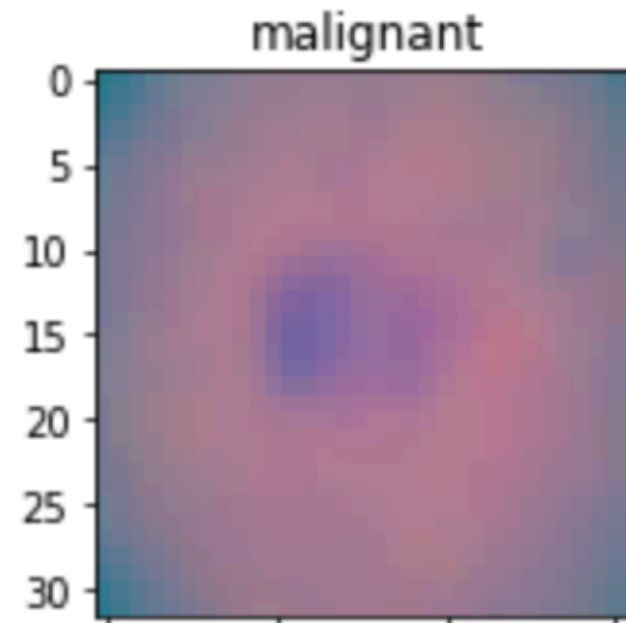
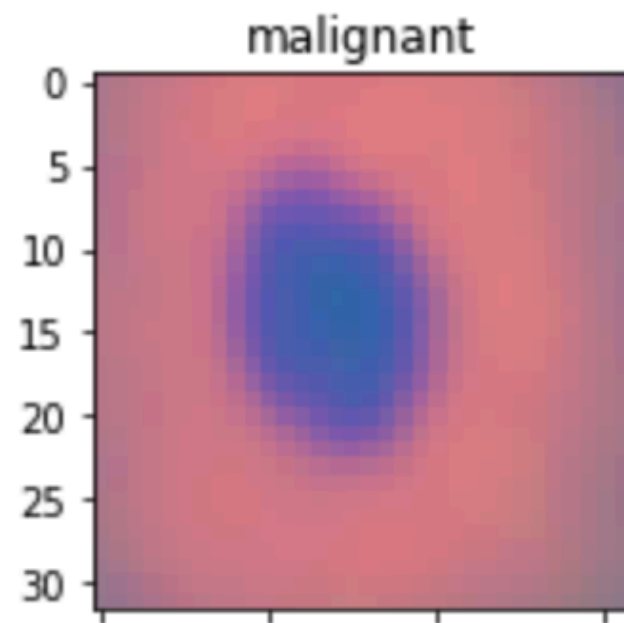


Image Preprocessing

- **Image scaling:-** resize the images to have fixed height and width
- **Color space transformation:-** RGB, grayscale
- **Contrast enhancement through histogram equalization, adaptive histogram equalization**
- **Morphology methods:-** Gaussian smoothing, threshold, erosion, dilation

Image preprocessing

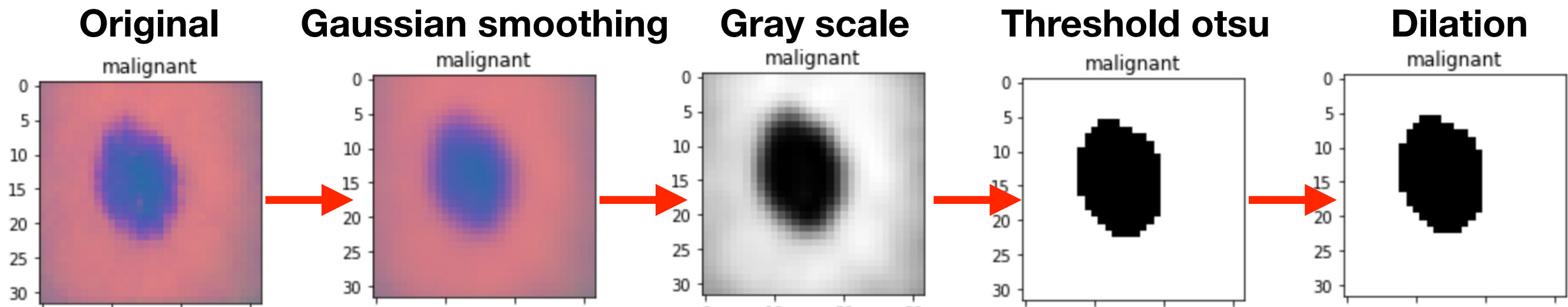
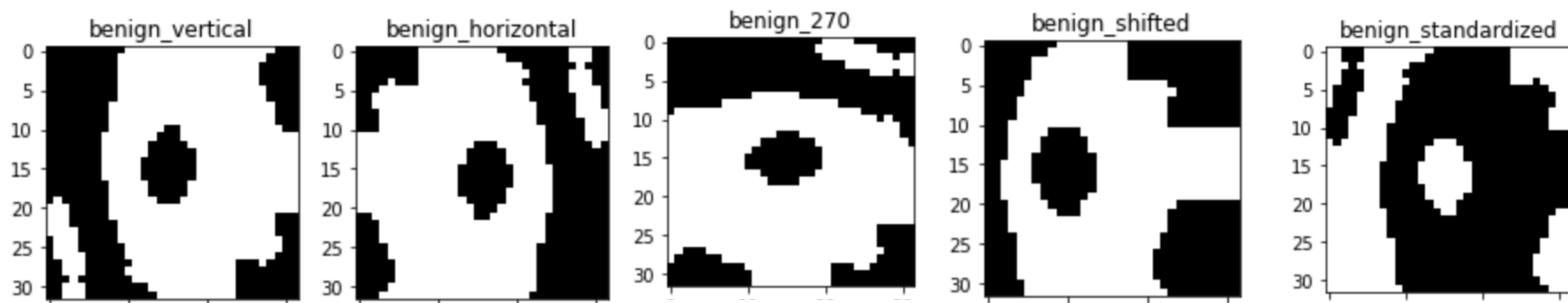


Image augmentation

- Random flip:- Horizontal or Vertical
- Random rotation
- Random shift, shear, zoom
- Random color jitter (brightness, standardize)

Augmented images in grayscale

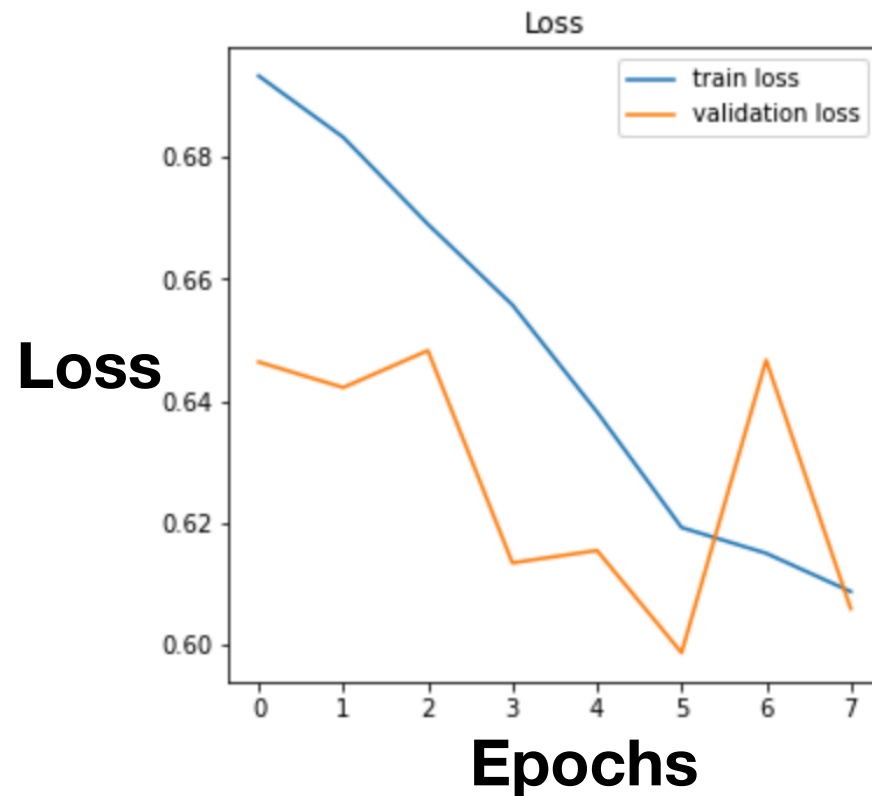


Modeling

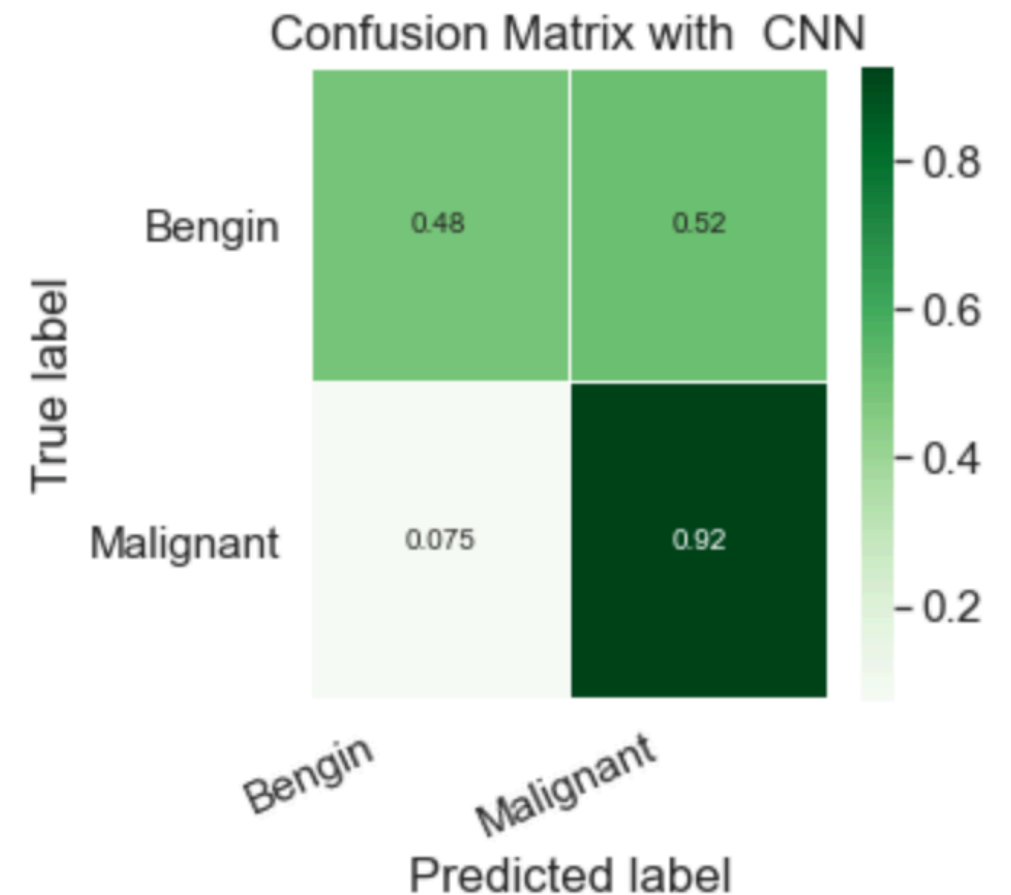
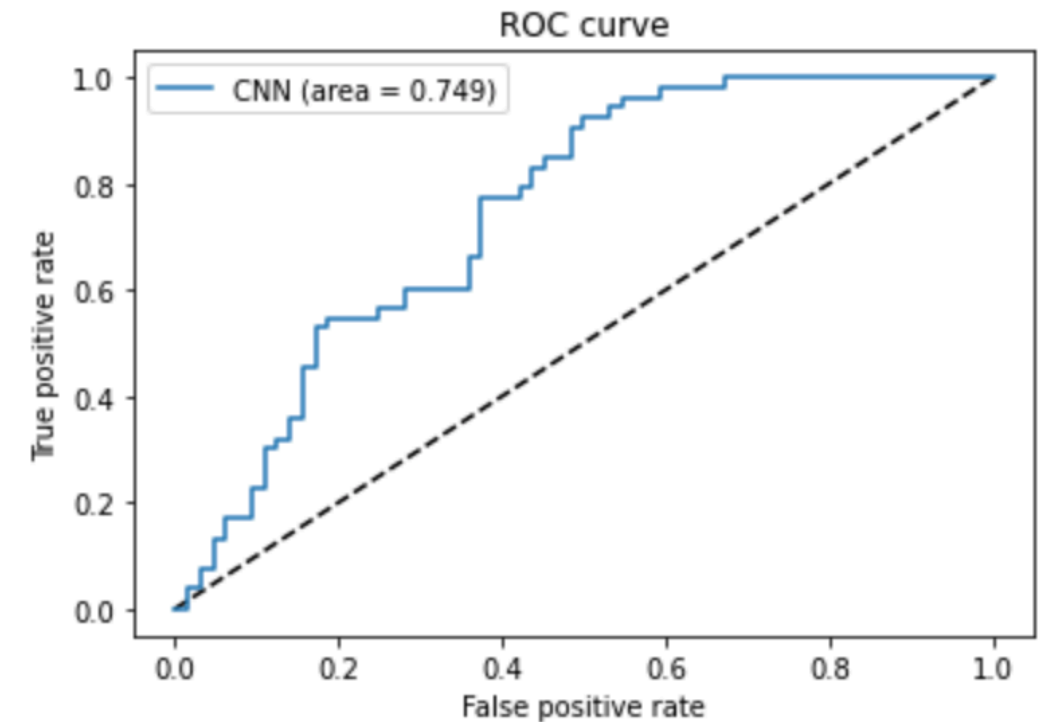
- Built model on equalized dataset
- Image normalization with pixel value between 0 and 1, RGB images with 3 channels
- Deep neural network: four convolutional layers, four max-pooling layers and two dense layers
- Convolutional layers:-
 - First layer: 32-3 x 3 filters
 - Second layer: 64-3 x 3 filters
 - Third layer:- 128-3 x 3 filters
 - Fourth layer- 128-3 x 3 filters
- Max-pooling layers each of the size 2x2 with strides 2
- Optimizer: Adam
- Loss function: Binary crossentropy
- Metrics: AUC
- Regularization with image augmentation, dropout and early stopping

Model evaluation

Loss function with training epochs



- Model predicts malignant images with a probability of 0.92
- Model critical in predicting malignant images than benign images
- The cost of incorrectly classifying malignant images are much riskier than incorrectly classifying benign images

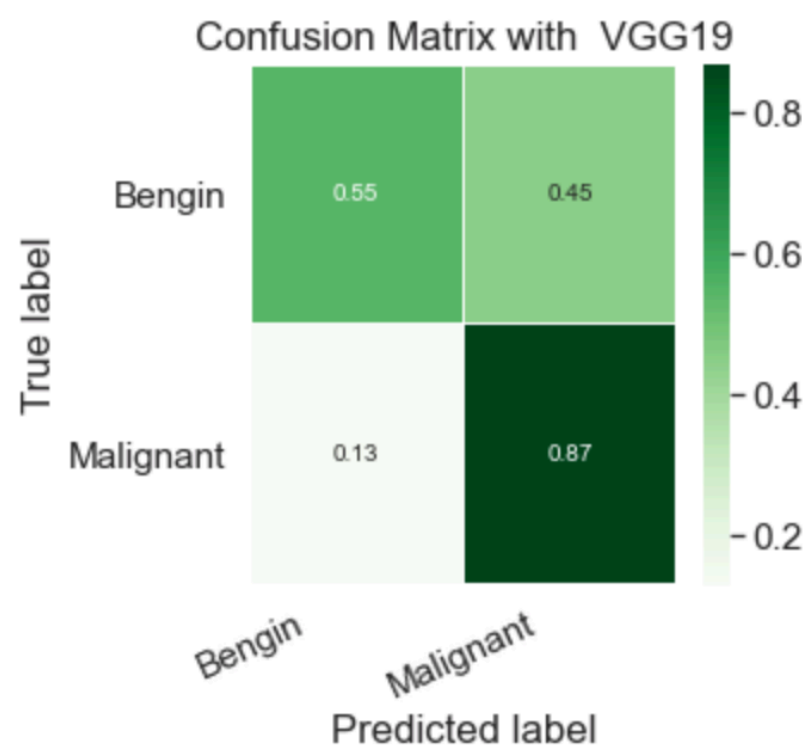
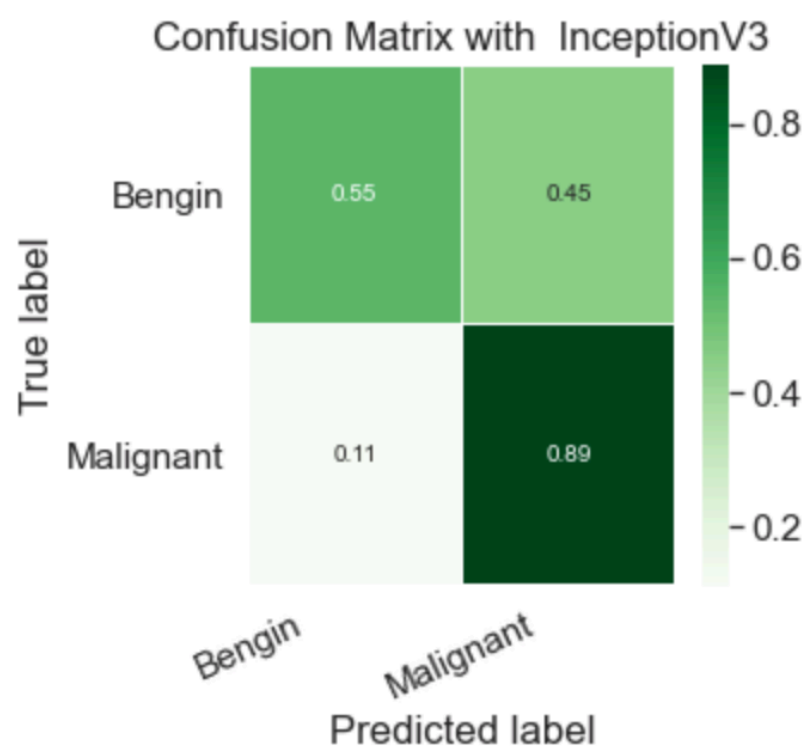
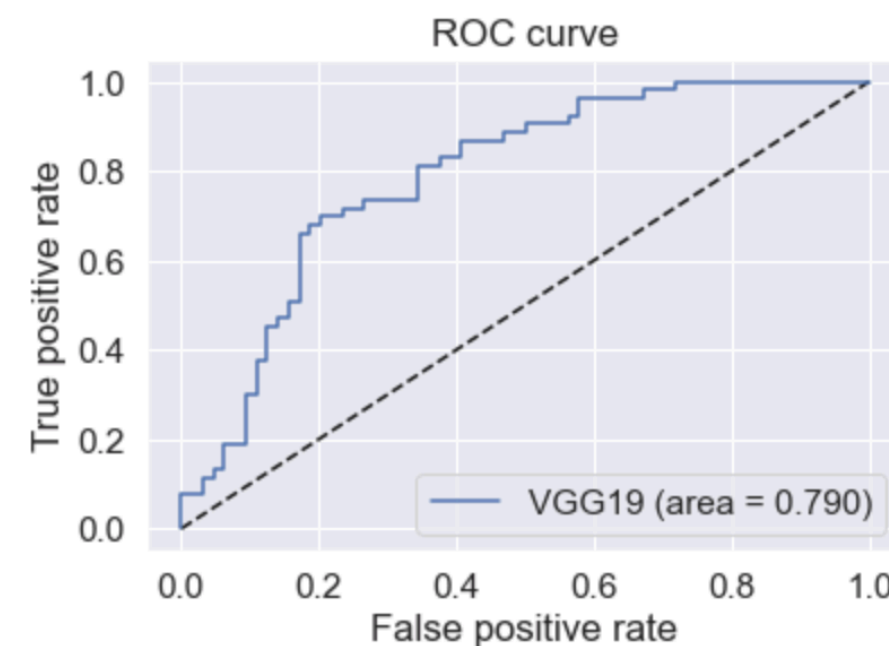
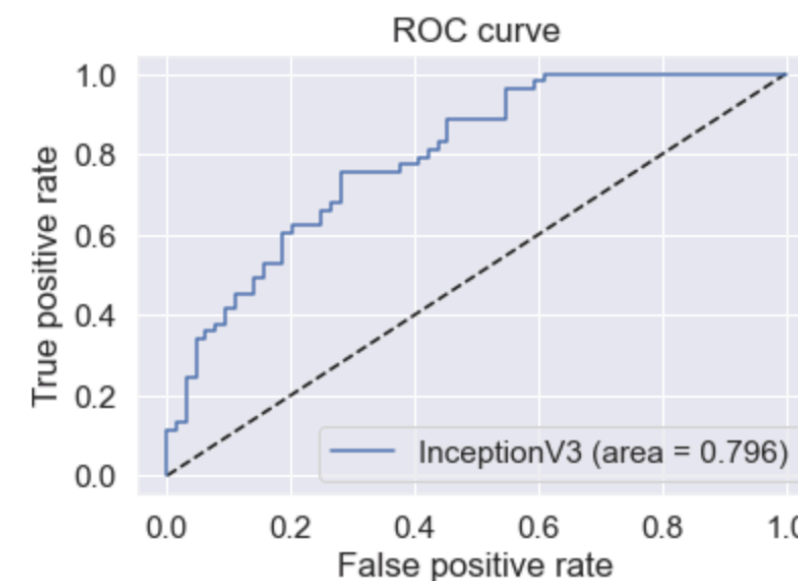


Transfer learning

- **Build models with weights trained on imageNet and EfficientNet**
- **Start the learning process from the patterns that have already been learned**
- **Employed models**
 - **VGG19**
 - **Resnet50 and Resnet152**
 - **InceptionV3**
 - **EfficientNetB0 and EfficientNetB5**
- **Trained model fixing all layers or allowing few top layers to be trained**

Model evaluation

| Model | Loss | AUC | Precision | Recall | F-score |
|--------------------|-------|------|-----------|--------|---------|
| CNN | 0.59 | 0.75 | 0.48 | 0.92 | 0.63 |
| VGG19 | 0.63 | 0.79 | 0.55 | 0.87 | 0.67 |
| Resnet50 and 152 | 0.71 | 0.70 | 0 | 1 | 0 |
| Inception V3 | 0.60 | 0.80 | 0.55 | 0.89 | 0.68 |
| EfficientNetB0 and | 0.695 | 0.39 | 0 | 1 | 0 |



Conclusions

- **Developed binary models that can identify images to a benign or malignant class**
- **Multiple models were built using deep neural networks and transfer learning architecture**
- **CNN achieved precision of 0.48, recall of 0.92, F1 score of 0.63 and AUC of 0.75**
- **VGG19 and inceptionV3 model performed better than CNN on precision, F1 score and AUC**
- **Resnet and EfficientNet couldn't performed better than CNN**

Recommendations

- **Develop model on imbalanced dataset**
- **Fine tune CNN with hyper parameters such as batch size, learning rate, average pooling, filter size and strides**
- **Evaluate model performance with other pre-trained models such as Xception, VGG16, MobileNet, Densenet, NASNet, MobileNetV2**
- **Fine tune pre-trained models freezing all layers or making few layers trainable**