# SKIN CANCER IMAGE CLASSIFICATION WITH CONVOLUTIONAL NEURAL NETWORKS

#### **Problem Statement**

- Considering the limited availability of the resources, early detection of skin cancer is highly important.
- Accurate diagnosis and feasibility of detection are vital in general for skin cancer prevention policy.
- Skin cancer detection in early phases is a challenge for even the dermatologist.

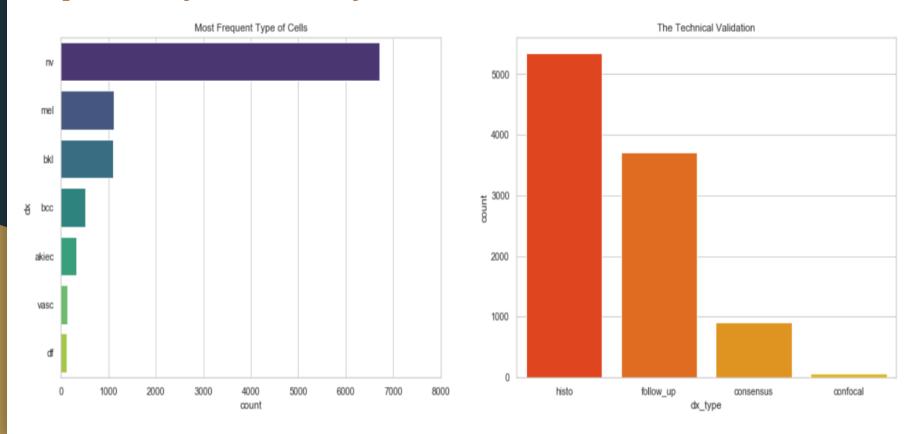
## Objective

- To detect 7 different classes of skin cancer using Convolution Neural Network with keras tensorflow in backend.
- To analyse the result to see how the model can be useful in practical scenario.
- Diagnosing methodology using Image Processing and Deep Learning models.

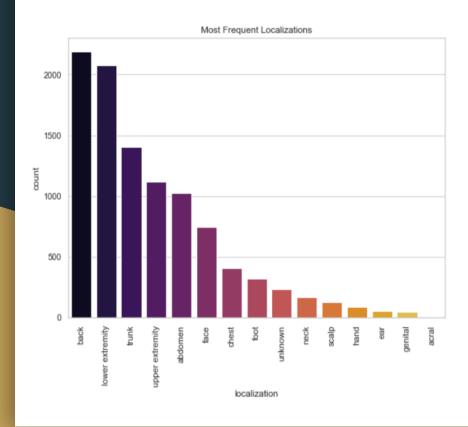
## Data Description

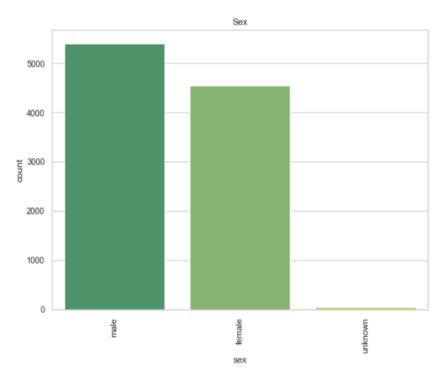
- The dataset that we are using is the HAM10000 dataset.
- 10015 dermatoscopic images.
- 7 features lesion id, image id, dx, dx\_type, age, sex, localization
- It has 7 different classes of skin cancer:
  - 1. Melanocytic nevi
  - 2. Melanoma
  - 3. Benign keratosis
  - 4. Basal cell carcinoma
  - 5. Actinic keratoses
  - 6. Vascular lesions
  - 7. Dermatofibroma

#### **Exploratory Data Analysis** - Type of Skin Cancer and Technical Validation



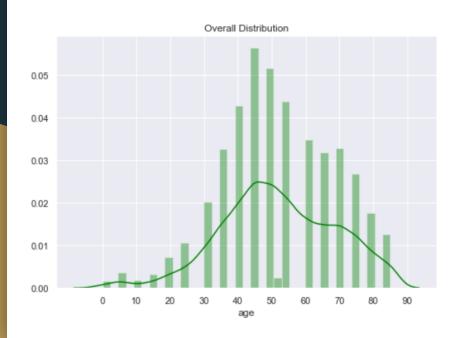
#### **Exploratory Data Analysis - Localization and Sex**

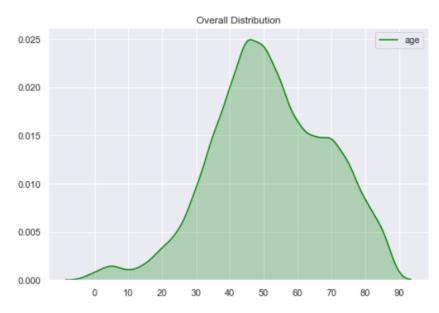




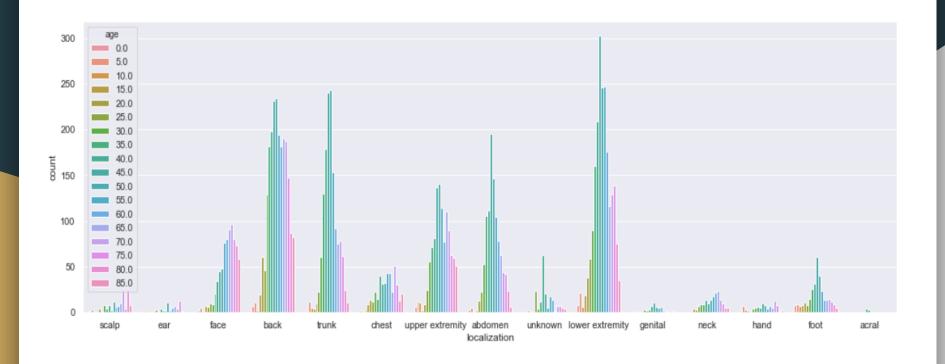
#### Exploratory Data Analysis - Age

#### Distribution of Age

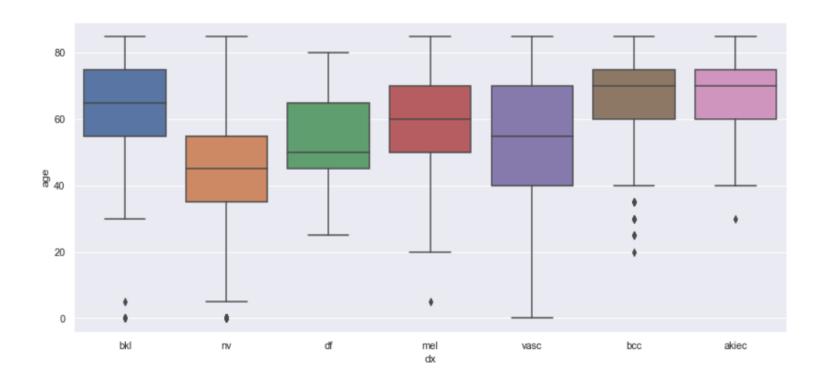




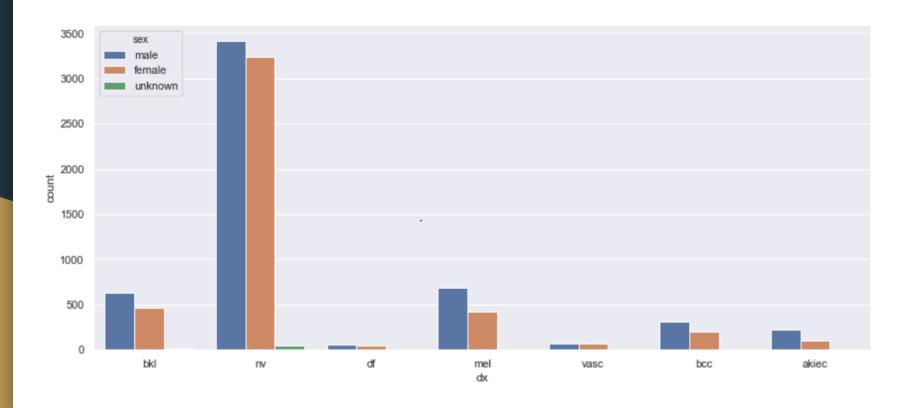
#### **Exploratory Data Analysis - Correlation between Localization and Age**



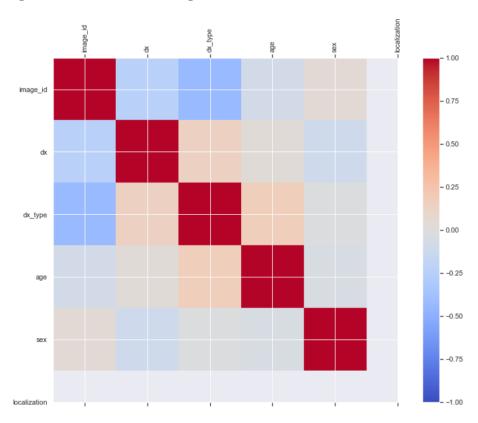
#### **Exploratory Data Analysis** - Correlation between Age and Type of Cancer



#### **Exploratory Data Analysis - Correlation between Type of Cancer and Sex**



#### **Exploratory Data Analysis** - Correlation between parameters

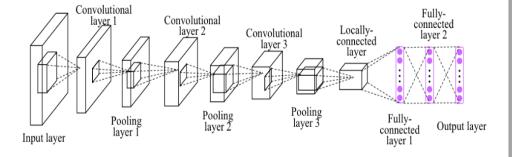


## Data Preprocessing

- Remove the Null Values
- Remove the Duplicates
- Splitting the dataset into Training and Testing sets
- Convert categorical columns into Numerical columns using One Hot Encoding and Label Encoding
- Resizing the Images
- Normalization

#### **CNN** Architecture

- 1. Convolution layer -Conv2D
- 2. Pooling layer MaxPooling2D
- 3. Flatten layer
- 4. Fully connected layer -Dense



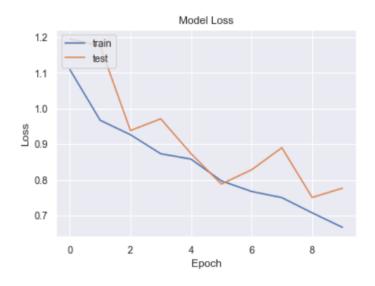
#### **CNN** Architecture

```
model = Sequential()
model.add(Conv2D(32, kernel size=(3, 3),activation='relu',input shape=(100,100,3),padding='same'))
model.add(MaxPooling2D((2, 2),padding='same'))
model.add(Dropout(0.20))
model.add(Conv2D(64, (3, 3), activation='relu',padding='same'))
model.add(MaxPooling2D(pool size=(2, 2),padding='same'))
model.add(Dropout(0.40))
model.add(Conv2D(128, (3, 3), activation='relu',padding='same'))
model.add(LeakyReLU(alpha=0.1))
model.add(MaxPooling2D(pool size=(2, 2),padding='same'))
model.add(Dropout(0.20))
model.add(Flatten())
model.add(Dense(64, activation='linear'))
model.add(LeakyReLU(alpha=0.1))
model.add(Dense(128, activation='linear'))
model.add(Dense(256, activation='linear'))
model.add(Dense(7, activation='softmax'))
model.summary()
```

## Setting Optimizer and Fitting the model

```
: model.compile(optimizer='adam',loss='categorical crossentropy',metrics=['accuracy'])
history=model.fit(train X,one hot train,batch size=128,epochs=10,validation split=0.2)
Train on 6409 samples, validate on 1603 samples
Epoch 1/10
v: 0.6569
Epoch 2/10
v: 0.6438
Epoch 3/10
v: 0.6644
Epoch 4/10
v: 0.6837
Epoch 5/10
v: 0.6893
Epoch 6/10
y: 0.7137
Epoch 7/10
v: 0.6993
Epoch 8/10
v: 0.6949
Epoch 9/10
v: 0.7118
Epoch 10/10
y: 0.7255
```

### **Model Evaluation**





#### Conclusion

Accuracy is higher if model is trained on more samples of lower resolution than small samples of high resolutions. We have achieved the accuracy of 75%.

## Future Scope

Going forward, we can continue to refine the model to achieve a stable decrease in loss function with every epoch, build an interface such that given an image of a skin lesion within the two classes, the output will give a % probability of which of the seven classes it belongs to.

Q & A

Thank You