Melanoma Detection - Case Study

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Overview

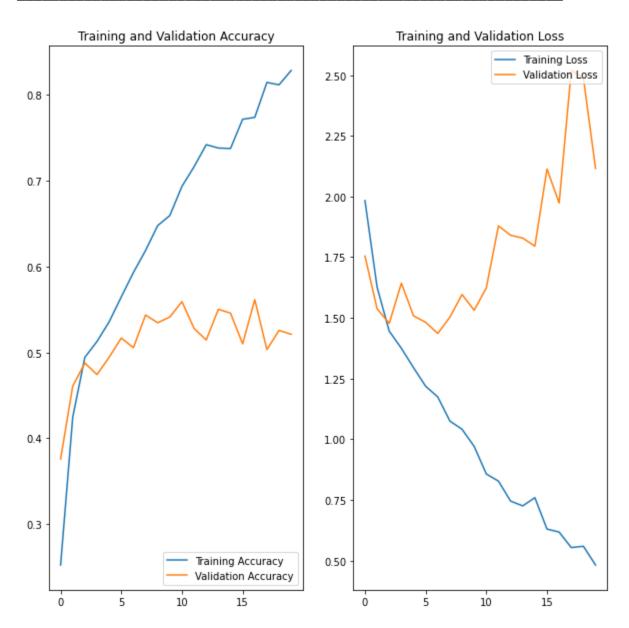
Basic Model

For the basic model, the first layer in the model is a Rescaling layer that rescales the image by 255. We use the original dataset for this model. I am using a 3 CNN unit structure. All CNN Layers have 1 Conv2D layer followed by MaxPooling2D layer. The CNN units have an increasing number of filters going from 32 to 128. After the CNN units, we have a flatten layer and then a Dense layer with 9 (number of classes) layers.

Model: "sequential"

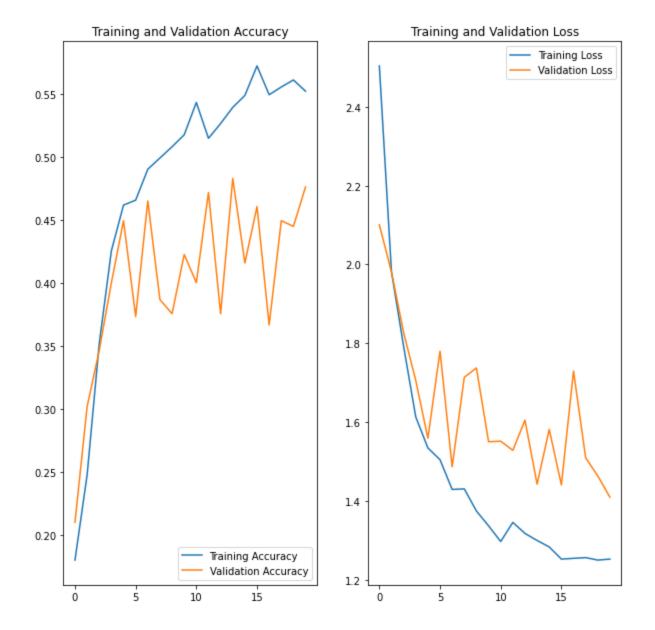
| | Output Shape | Param # |
|--|----------------------|---------|
| rescaling (Rescaling) | | 0 |
| conv2d (Conv2D) | (None, 180, 180, 32) | 896 |
| <pre>max_pooling2d (MaxPooling2D)</pre> | (None, 90, 90, 32) | 0 |
| conv2d_1 (Conv2D) | (None, 90, 90, 64) | 18496 |
| <pre>max_pooling2d_1 (MaxPooling 2D)</pre> | (None, 45, 45, 64) | 0 |
| conv2d_2 (Conv2D) | (None, 45, 45, 128) | 73856 |
| <pre>max_pooling2d_2 (MaxPooling 2D)</pre> | (None, 22, 22, 128) | 0 |
| flatten (Flatten) | (None, 61952) | 0 |
| dense (Dense) | (None, 9) | 557577 |

Total params: 650,825
Trainable params: 650,825



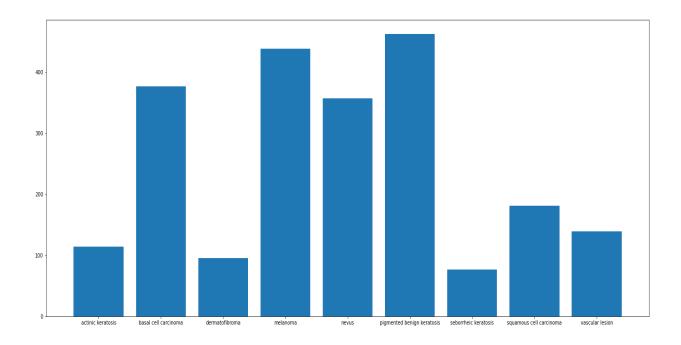
Data Augmentation and Drop Outs

In this case, we update the basic model and add dropouts after each CNN Unit. We can see adding dropout layers reduces the over fitting issue.



Distribution in the Dataset

We can visualise the initial distribution of the dataset and observe that there is data imbalance between the classes. It would be a good idea to augment the data so that we reduce the sparsity of the dataset.



Data Augmentation

We add 500 augmented images to each class to reduce the sparsity of the classes with lower numbers of data. The updated number of images in each class is as shown below:

| pigmented benign keratosis | 962 |
|----------------------------|-----|
| melanoma | 938 |
| basal cell carcinoma | 876 |
| nevus | 857 |
| squamous cell carcinoma | 681 |
| vascular lesion | 639 |
| actinic keratosis | 614 |
| dermatofibroma | 595 |
| seborrheic keratosis | 577 |

BatchNormalisation

We add a batch normalisation layer to improve the model performance.

Model: "sequential_3"

| Layer (type) | Output | Shape | e | | Param | # |
|---|--------|-------|-------|---------|-------|---|
| ======================================= | | | | ======= | | |
| rescaling_2 (Rescaling) | (None, | 180, | 180, | 3) | 0 | |

| <pre>batch_normalization (BatchN ormalization)</pre> | (None, 180, 180, 3) | 12 |
|--|----------------------|--------|
| conv2d_6 (Conv2D) | (None, 180, 180, 32) | 896 |
| <pre>batch_normalization_1 (Batc hNormalization)</pre> | (None, 180, 180, 32) | 128 |
| <pre>max_pooling2d_6 (MaxPooling 2D)</pre> | (None, 90, 90, 32) | 0 |
| dropout_3 (Dropout) | (None, 90, 90, 32) | 0 |
| conv2d_7 (Conv2D) | (None, 90, 90, 64) | 18496 |
| <pre>max_pooling2d_7 (MaxPooling 2D)</pre> | (None, 45, 45, 64) | 0 |
| dropout_4 (Dropout) | (None, 45, 45, 64) | 0 |
| conv2d_8 (Conv2D) | (None, 45, 45, 128) | 73856 |
| <pre>max_pooling2d_8 (MaxPooling 2D)</pre> | (None, 22, 22, 128) | 0 |
| dropout_5 (Dropout) | (None, 22, 22, 128) | 0 |
| flatten_2 (Flatten) | (None, 61952) | 0 |
| dense_2 (Dense) | (None, 9) | 557577 |

Total params: 650,965 Trainable params: 650,895 Non-trainable params: 70

