

Configuration Manual

MSc Research Project Data Analytics

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Configuration Manual

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1 Introduction

The specifications for the A Comparative Study of Pixel Values and Landmark Detection Features to Solve for Facial Emotion Recognition research are described in this configuration manual. It covers hardware and system specifications that should be considered minimum requirements for the study results replication. The following sections cover all phases of the implementation process and the overall assessment of the research work.

2 System Specification

2.1 Configuration of Hardware

A M1 MacBook pro 8GB was used for the entire study. Figure 1 shows the specific information regarding Device, Chip, Memory and OS also listed below.

- Device: MacBook Pro (13-inch, M1, 2020)
- Chip: Apple M1
- Memory: 8 GB



Figure 1: Device Specification

2.2 Software Configuration

Google Colaboratory (Colab) was used for the entire duration of the study. Due to the Mac M1 have a very laborious process to install TensorFlow which can be easily accessed by utilizing Colab online. This meant that a combination of Google Drive and Google Colab had to be used to importing the datasets used. Figure 2 displays the landing page for google colab¹

Software Requirements below:

- **OS**: macOS Monterey v 12.5
- Google Account to access Google Drive to store files.
- Google Basic Subscription Plan for 100mb of drive storage (some datasets were over the free storage capacity.).
- Same google account can be used to access Google Colab.
- Google Colab Pro+ subscription (enables access to higher RAM and GPU, TPU.).

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	CO Markdow	n Guide				an
	CO Charts in	Colaboratory				I GP
	CO External	data: Drive, Sheets, a	nd Cloud Storage			
	CO Getting s	tarted with BigQuery				
						enu,
<>	_	_	_	New notes	oook Cancel	
	- More mei	mory				
						• ×

Google Colab

Figure 2: Colab Landing Page

As features extracted were numerous, particularly for 224 x 224 pixel images in datasets that were augmented resulting in dataframes with large number of columns memory crashes. At times the data was over 35k rows and 150k columns. This mostly happened during the experimentation phase of the project and it is not part of the needed steps to complete the study but as RAM was crashing at around 25gb+ a decision was reached to upgrade to Colab Pro+ which unlocked 54gb of RAM as per Figure 3.

GPU was also available and used during the CNN creation. Figure 4 shows the configuration in Google Colab. Figure 5 displays how to confirm that GPU is available and how to select High RAM.

¹Google Colaboratory: https://colab.research.google.com/

Google Colab Memory



You are using a high-RAM runtime!

Figure 3: Colab Memory





Figure 4: Selecting GPU

Google Colab GPU

0	<pre>1 ## checking if connected to GPU for faster times 2 3 gpu_info = lnvidia-smi 4 gpu_info = '\n'.join(gpu_info) 5 if gpu_info.find('failed') >= 0: 6 print('Not connected to a GPU') 7 else: 8 print(gpu_info)</pre>	
C⇒	Wed Aug 10 22:54:19 2022 	+
	GPU Name Persistence-M Bus-Id Disp.A Volatile Uncorr.F Fan Temp Perf Pwr:Usage/Cap Memory-Usage GPU-Util Compute I I Image: March and a straight and straight and straight and a strain and a straight and straight a	CC M. M.
	0 Tesla P100-PCIE Off 0000000:00:04.0 Off	0 1t /A
		+
	GPU GI CI PID Type Process name GPU Memo ID ID Usage	ry



3 Database Sources

Many different datasets were requested to be part of this study. Due to time limitations and resources constraints two datasets were selected to the final report.

- 1. The posed dataset Japanese Female Facial Expression (JAFFE) created by Lyons et al. consists of 10 Japanese female subjects displaying the 6 basic facial emotions plus neutral emotion. There are 213 images in total in 256 by 256 pixes grayscale. Images are in the Tiff format with no compression. Images were created with excellent lighting and background. This dataset has over 25 years and is utilized in many different papers (Lyons; 2021).
- 2. The unposed dataset Static Facial Expressions in the Wild (SFEW) (Dhall et al.; 2012) was selected to contrast the JAFFE dateset for having "in-the-wild" type of images. The dataset contains 700 images and 68 subjects. The images were collected from different movies and display a variety of genders, occlusions, different illumination and poses, being referred as close to real life environment by the authors.

4 Libraries Used, Image Importing and Train/Test Split

This section looks into libraries used and how to point to the right directory and import the images and extract Features. The code is the same for both datasets the only thing that changes is the directory pointing at either JAFFE or SFEW folders.

4.1 Libraries Used

Figure 6 shows all the used libraries to execute the code. Please note that Augmentor library is commented out as it required to be "!pip installed" every time the google colab notebook was switched between GPU and normal RAM. Once the process of Data Augmentation was completed with Augmentor it was commented out to avoid this step.

4.2 Loading the data

The first step after loading the libraries is to point at where the folder with the dataset chosen is. Figure 7 displays the code.

4.3 Data Exploration and First Look at Landmarks

The following code is a loop that goes into each of the emotion folders, grabs one image, converts from BGR to RGB (more details in the paper), resizes it to 224 x 224 pixels, labels it based on the name of the folder. Extract 68 landmark features from the face and displays both the resulting image created from the points and the superimposed image above the original face. The code then breaks and jumps into the next folder until all named folders have been "visited". Figure 8 shows the code utilized and Figure 9 some of the resulting emotions and faces.

Imported Libraries



Figure 6: All Imported Libraries

Path and Emotion Folder Names

[] 1 PATH = "/content/drive/MyDrive/Masters/jaffedbase manual" 2 CATEGORIES = 'Angry', 'Disgust', 'Fear', 'Happy', 'Neutral', 'Sad', 'Surprise' ### 0, 1, 2, 3, 4, 5, 6

Figure 7: Pointing to the Right Path

Emotion Labels and Landmarks Code

0	1 de	f show_landmarks(PATH):
-	2	landmarks = []
	3	for x in CATEGORIES:
	4	<pre>path = os.path.join(PATH, x)</pre>
	5	<pre>for img in os.listdir(path):</pre>
	6	<pre>img_array = cv2.imread(os.path.join(path, img))</pre>
	7	<pre>img_array = cv2.cvtColor(img_array, cv2.COLOR_BGR2RGB)</pre>
	8	<pre>img_new = cv2.resize(img_array, (224, 224))</pre>
	9	<pre>landmarks = extract_face_landmarks(img_new)</pre>
	10	<pre>fig = plt.figure(figsize=(15, 5))</pre>
	11	$ax = fig.add_subplot(1, 3, 1)$
	12	<pre>plt.title(f'Emotion: {x}')</pre>
	13	ax.imshow(img_new)
	14	<pre>ax = fig.add_subplot(1, 3, 2)</pre>
	15	<pre>ax.scatter(landmarks[:, 0], -landmarks[:, 1], alpha=0.8)</pre>
	16	<pre>ax = fig.add_subplot(1, 3, 3)</pre>
	17	<pre>img2 = img_new.copy()</pre>
	18	
	19	for p in landmarks:
	20	<pre>img2[p[1]-3:p[1]+3, p[0]-3:p[0]+3, :] = (255, 255, 255)</pre>
	21	# note that the values -3 and +3 will make the landmarks
	22	# overlayed on the image 6 pixels wide; depending on the
	23	# resolution of the face image, you may want to change
	24	# this value
	25	
	26	<pre>ax.imshow(img2)</pre>
	27	<pre>plt.title(f'Emotion: {x}')</pre>
	28	plt.show()
	29	break

Figure 8: Emotion Labels and Landmarks Code



Emotion Labels and Landmarks Results

Figure 9: Emotion Labels and Landmarks Results

Variation of the code in figure 10 import the data either as Pixel Values, 68 Landmarks or the calculation of the euclidean distances between landmark points. Figure 10 shows a version of the code and the main difference is explained in the list below.

read data Function



Figure 10: read data Function

The version of the code with the two lines below after image is resized with img_new = cv2.resize(img_array, (IMGSIZE, IMGSIZE)) collects flattened RGB pixel values.

- img_new = np.array(img_new).flatten() (flattens the np array image after it was resized).
- data.append([img_new, class_num]) (appends the information that was just flattened to the data empty list).

The following 3 lines instead of the ones above after image resize takes place (img_new = cv2.resize(img_array, (IMGSIZE, IMGSIZE))) collects information on the 68 landamrk points.

- landmarks = extract_face_landmarks(img_new) (extracts 68 facial landmark points using the Mlxtend library).
- landmarks = np.array(landmarks).flatten() (*flattens the landmark information*).
- data.append([landmarks, class_num]) (appends the information that was just flattened to the data empty list).

The following 3 lines instead of the ones above after image resize takes place (img_new = cv2.resize(img_array, (IMGSIZE, IMGSIZE))) collects information on the 68 landamrk points.

- landmarks = extract_face_landmarks(img_new) (extracts 68 facial landmark points using the Mlxtend library).
- distances = compute_landmark_distances(landmarks) (utilizes the code function to calculate euclidean distance).
- data.append([distances, class_num]) (appends the information that was just flattened to the data empty list).

Function to calculate euclidean distances



Figure 11: Function to calculate euclidean distances

Additional function to calculate euclidean distance between points can be seen in figure 11.

Once data is read, it needs to be shuffled, random shuffle is used, first setting the seed so results can be replicated, second shuffling the data and finally assigning the data to X and y variables and changing from list to np.array. For pixel values, the values can be normalised by dividing X = X/255.0. After everything is ready a pickle file is created so that in future data reading step doesn't have to be repeated. Figure 12 shows it in details.

reading, shuffling and saving pickle file

Figure 12: reading, shuffling and saving pickle file

4.4 Looking for missing values

Transforming the data from np.array to dataframe allows us to look into the data better and see if any values are null or 0.0000 where landmark detection could not read faces. This didn't happen for JAFFE dataset or for extraction of pixel values but happened for SFEW over 26k times with data augmented and normal dataset. Figure 13 shows how the table looks like when there's no missing data. Figure 14 shows the table with missing data.

There's no need to try to trust your vision only as a simple code df = df.loc[(df.iloc[:, 0:4623] == 0).all(axis=1)] removes the missing values and the table is ready to be saved

No missing data

1	1 # 2 3 d 4 d	## CI t = 1 t('0)	becking f pd.DataFr riginal_1	or mission ame(X) abel'] =	y values														
1	1 d	ť																	
		۰	1	2	3	4	5	6	7	8	,	 4615	4616	4617	4618	4619	4620	4621	4
	0	0.0	0.250555	0.501110	0.769560	1.013794	1.229386	1.426924	1.641476	1.794126	1.890914	0.190029	0.235702	0.0000000	0.100000	0.183333	0.333750	0.183333	0.100
	1	0.0	0.250565	0.520160	0.758637	1.000278	1.224972	1.461069	1.677051	1.830604	1.962562	 0.177169	0.165867	0.066667	0.120185	0.195078	0.366667	0.200000	0.10
	2	0.0	0.250565	0.485913	0.729610	0.954521	1.181336	1.377296	1.557954	1.719677	1.833106	0.166667	0.184089	0.037268	0.105409	0.186339	0.417000	0.200000	0.10
	3	0.0	0.235702	0.454911	0.693221	0.905692	1.122621	1.313181	1.515292	1.666667	1.791957	 0.142400	0.216667	0.133333	0.177169	0.263523	0.380068	0.217307	0.11
	4	0.0	0.233928	0.469338	0.709555	0.941925	1.170233	1.389744	1.580436	1.728840	1.807776	 0.134371	0.190029	0.184089	0.195078	0.250000	0.368932	0.166667	0.083
	206	0.0	0.267187	0.537484	0.791974	1.055804	1.291532	1.538488	1.774511	1.932255	2.044573	0.164992	0.153659	0.116867	0.165867	0.225691	0.372678	0.184089	0.10
	207	0.0	0.300000	0.603592	0.895979	1.184389	1.465550	1.767374	2.005688	2.158253	2.230159	 0.169967	0.5353333	0.400347	0.416667	0.440328	0.471699	0.216687	0.113
	208	0.0	0.217307	0.438432	0.660387	0.889444	1.085766	1.284631	1.468654	1.608657	1.724658	0.120185	0.174005	0.065667	0.097183	0.179505	0.403113	0.183333	0.083
	209	0.0	0.250555	0.504425	0.759020	0.987140	1.213352	1.424001	1.595131	1.738454	1.848122	0.177169	0.150923	0.016667	0.083333	0.184089	0.334996	0.183333	0.083
	210	0.0	0.266667	0.533594	0.801561	1.058432	1.294433	1.539120	1.770515	1.924765	2.000462	0.188562	0.165657	0.065657	0.097183	0.195078	0.350000	0.183333	0.05
	11 15	x enc	4625 colum	19															

Figure 13: No missing data

Missing data

	٥	1	2	3	4	5	6	7		,	 4615	4616	4617	4618	4619	4620	462
0	0.0	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	 0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.00000
1	0.0	0.0000000	0.000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	 0.0000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.0000
2	0.0	0.0000000	0.000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.00000
з	0.0	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.0000
4	0.0	0.000000	0.000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.000000	0.000000	 0.0000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.00000
1389	0.0	0.117851	0.217307	0.320590	0.396162	0.479873	0.573973	0.643126	0.690008	0.719761	 0.074538	0.101379	0.016867	0.023570	0.062706	0.089753	0.05270
1390	0.0	0.000000	0.000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.000000	0.000000	 0.0000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.0000
1391	0.0	0.084984	0.183333	0.266667	0.351584	0.454911	0.529675	0.579751	0.626276	0.674949	0.060093	0.037268	0.0000000	0.037268	0.068718	0.153659	0.0687
1392	0.0	0.0000000	0.000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.0000
1393	0.0	0.0000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.000000	0.0000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.000000	0.00000

Figure 14: missing data

as X again now without unwanted 0 values. After all is confirmed and pickled file saved it can be loaded using the code displayed in figure 15.

Reading pickle file

Figure 15: reading pickle file

4.5 Creating Train and Test Split

The code in Figure 16 splits X and y into train and test data that can be used to train a model.

5 Training a ML model

This section follows the code in the colab book named "Machine Learning Techniques". Once data is assigned to the X_train and X_test, y_train and y_test it is ready to be used with the different ML models.

Baseline models are run as per Figure 17.

figure 18 shows an example of the grid search parameters.

Follow the code lines running both baseline models and grid search. Some of the grid search combination took above 11 hours to run.

Train and Test Split

ating X train/v train/X Test/v test

5100	Anny Action (Josef Jose
D	<pre>1 # create train/test split 2 X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=0.23, stratifyry, shuffle=True, random_state=42) 4 ### elited from code <u>https://www.theythoncode.com/article/building-a-speech-emotion-recognizer-using-shlearn 5 # details 7 # o samples in training data 8 print(' ' Sumber of training samples', X_train.shape(0)) 9 # o samples in training the testing data 9 # o samples in training the testing data 9 # o samples in training samples', X_train.shape(0)) 9 # o samples in training samples', X_train.shape(0)) 9 # o samples in testing data 9 # o samples i</u></pre>

Figure 16: Creating Train and Test Split

Baseline model

Figure 17: Baseline model

Grid search parameters

Figure 18: Grid search parameters

5.1 Data Transformation

Two types of data transformation were used to reduce dimensionality in the data LDA transformation and PCA. Once new X_t train and X_t test variables were created using PCA or LDA methods the same models were ran again, with the new variables, so that results could be compared.

6 Data Augmentation

Data augmentation was used to increase the number of image examples to train the model. Both a library called Augmentor and a Keras approach were used at different times in the process of the code creation. Augmentor generated images in a folder called output in the different emotion classes. These were then manually moved and reorganised but as Mlxtend library struggled to find faces in the data this step was not used in the main project data. Augmentor steps can be seen in figure 19.

Augmentor Data Augmentation

Data	Augmentation
0	<pre>1 p = Augmentor.Pipeline("/content/drive/MyDrive/Masters/Sfew/Train/Disgust") #disgust increasing from 66 to 5k examples</pre>
C+	Initialised with 66 image(s) found. Output directory set to /content/drive/HyDrive/Hasters/Sfew/Train/Disgust/output.
[]	<pre>1 p.random_distortion(probability=1, grid_width=2, grid_height=2, magnitude=4) 2 p.fitp_left_right(probability=1) 3 p.roste(probability=1, matheft_rostation=4, matheft_rostation=4) 4 p.orcop_random(probability=1, percentage_area=0.95)</pre>
[]	1 p.sample(5000)
	Processing <pil.image.image 0x7f08b03451390="" at="" image="" mode="RGB" size="684x347">: 100%</pil.image.image>
[]	1 p = Augmentor.Pipeline("/content/drive/MyDrive/Masters/Sfew/Train/Angry") #Angry increasing from 178 to 5k examples
	Initialised with 178 image(s) found. Dutput directory set to /content/drive/NyOrive/Masters/Sfew/Train/Angry/output.
[]	<pre>l p.random_distortion(probability=1, grid_width=2, grid_height=2, magnitude=4) 2 p.fitp_left_right(probability=1) 3 p.rostes(probability=1, grid_ft_rostation=4, max_right_rostation=4) 4 p.orcop_random(probability=1, percentage_area=0.95)</pre>
[]	1 p.sample(5000)
	Processing <pil.image.image 0x7f8b8bd59690="" at="" image="" mode="RGB" size="684x547">: 100%</pil.image.image>

Figure 19: Augmentor Data Augmentation

Figure 20 shows the data augmentation that was used in the creation of the CNN. Although results did not increase with the data augmentation so this step is not present in the main accuracy of the CNN model.

Figure 20: CNN Test Data

7 CNN Setup

To import data for the CNN with the right dimension the code is almost the same as the pixel value feature extraction code but without the .flatten() line as displayed by figure 21.

Importing data for CNN

Figure 21: Importing data for CNN

Run the code to create the function that plots the history of the model training as seen in figure 22. Followed by the code with the CNN layers in figure 23. and to compile the model in figure 24.

CNN History

-		
O	1 def	<pre>plot_history(history):</pre>
	2	<pre>fig, axs = plt.subplots(2)</pre>
	3	
	4	# create accuracy sublpot
	5	<pre>axs[0].plot(history.history["accuracy"], label="train accuracy")</pre>
	6	<pre>axs[0].plot(history.history["val_accuracy"], label="val accuracy")</pre>
	7	<pre>axs[0].set_ylabel("Accuracy")</pre>
	8	<pre>axs[0].legend(loc="lower right")</pre>
	9	<pre>axs[0].set_title("Accuracy eval")</pre>
	10	
	11	# create error sublpot
	12	<pre>axs[1].plot(history.history["loss"], label="train error")</pre>
	13	<pre>axs[1].plot(history.history["val_loss"], label="val error")</pre>
	14	<pre>axs[1].set_ylabel("Error")</pre>
	15	<pre>axs[1].set_xlabel("Epoch")</pre>
	16	<pre>axs[1].legend(loc="upper right")</pre>
	17	<pre>axs[1].set_title("Error eval")</pre>
	18	
	19	plt.show()

Figure 22: CNN history

The next lines of code create the prediction function that will use the test data to test the model with data that hasn't been seen by the model yet. It runs the entire test folder through the model and lets you pick a random image to test individually. As seen in figure 25.

8 Model Ensemble

A model ensemble made of the best performing models of this study was created in an attempt to increase the maximum achieved accuracy. This ensemble was successful and increased accuracy from 87% to 89%. Image 26 shows the details of the ensemble.

CNN Layers

1 model = tf.kera	<pre>is.models.Sequential([#data_augmentation,</pre>
2	<pre>tf.keras.layers.Conv2D(16,(3,3), 1, activation='relu',input_shape = (224,224,3)), ### 1st conv layer</pre>
3	tf.keras.layers.MaxPool2D(),
4	
5	## 2nd conv layer
6	<pre>tf.keras.layers.Conv2D(32,(3,3), 1, activation='relu'),</pre>
7	tf.keras.layers.MaxPool2D(),
8	tf.keras.layers.BatchNormalization(),
9	
10	## 5th conv layer
11	tf.keras.layers.Conv2D(16,(3,3), 1, activation='relu'),
12	tf.keras.layers.MaxPool2D(),
13	keras.layers.Dropout(0.2),
14	tf.keras.layers.BatchNormalization(),
15	
16	### flatten output and feed it into dense layer
17	tf.keras.layers.Flatten(),
18	<pre>tf.keras.layers.Dense(512, activation='relu'),</pre>
19	
20	## output layer
21	<pre>tf.keras.layers.Dense(7, activation='softmax')</pre>
22	1)

Figure 23: CNN Layers

CNN Compiler

0	<pre>1 # comple model 2 # optimiser = keras.optim 3 4 # nodel.compile(optimiser 5 # notrions) 7 # peint("Creating EarlySt 8 # early_stopping_callback 5 nodel.compile(optimiser=k 10 model.compile(optimiser=k 11 andel.summary()</pre>	<pre>izers.Adam(learning_rate- moptimiser, enorical_crossentropy', 'accuracy']) opping Callback') = EarlyStopping(sonitor= erss.optimizers.Adam(lear EarlyStopping(sonitor='v</pre>).00)) Vallos'.patience*)) sing_reted.(00)).loss = tf.keras.losses.fparseCategoricalCrossentropy(from_logits=%sins), metrics=['accuracy'] 	
Þ	Nodel: "sequential_23"			
	Layer (type)	Output Shape	Paran #	
	conv2d_45 (Conv2D)	(None, 222, 222, 16)	448	
	<pre>max_pooling2d_45 (MaxPoolin g2D)</pre>	(None, 111, 111, 16)	0	
	conv2d_46 (Conv2D)	(None, 109, 109, 32)	4640	
	<pre>max_pooling2d_46 (MaxPoolin g2D)</pre>	(None, 54, 54, 32)	0	
	<pre>batch_normalization_22 (Bat chNormalization)</pre>	(None, 54, 54, 32)	128	
	conv2d_47 (Conv2D)	(None, 52, 52, 16)	4624	
	<pre>max_pooling2d_47 (MaxPoolin g2D)</pre>	(None, 26, 26, 16)	0	
	dropout_18 (Dropout)	(None, 26, 26, 16)	0	
	batch_normalization_23 (Bat chNormalization)	(None, 26, 26, 16)	54	
	flatten_15 (Flatten)	(None, 10816)	0	
	dense_34 (Dense)	(None, 512)	5538304	
	dense_35 (Dense)	(None, 7)	3591	
	Total params: 5,551,799 Trainable params: 5,551,703 Non-trainable params: 96			

Figure 24: CNN Compiler

CNN Test Data

[]	<pre>1 def predict(model, X, y): 2 # add a dimension to input data for sample - model.predict() expects a 4d array in this case</pre>			
	X = X[np.newaxis,] # array shape (new dimention, 130, 13, 1)			
	<pre>4 # perform prediction 6 prediction = model.predict(X) 7 8 # get index with max value 9 predicted index = np.argmax(prediction, axis=1)</pre>			
	<pre>10 11 print("Target: {}, Predicted label: {}".format(y, predicted_index))</pre>			
0	<pre> 1 ### here we evaluate the model using the test dataset, 2 ###this files have not been seen by the DL model 3 4 test_loss, test_acc = model.evaluate(X_test, y_test, verbose=2) 5 print('\nTest accuracy:', test_acc) 6 7 # pick a sample to predict from the test set (30 chosen at random) 8 X_to_predict = X_test[5] 9 y_to_predict = y_test[5] 10 11 # predict sample 12 predict(model, X_to_predict, y_to_predict) </pre>			
C⇒	2/2 - 0s - loss: 0.6847 - accuracy: 0.8140 - 26ms/epoch - 13ms/step			
	Test accuracy: 0.8139534592628479 Target: 2, Predicted label: [2]			
r 1	1			

Figure 25: CNN Test Data

Model Ensemble

Figure 26: Model Ensemble

The ensemble used a repetition of the same model 3 times which increased the accuracy rather than using the algorithm just once. each repeated model builds up on each other and using hard voting helps increase accuracy.

9 Confusion Matrix

Confusion matrix is used to display model accuracy and F1-score information. This is shown in detail per emotional class, support and macro weighted average.

References

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Confusion Matrix

Figure 27: Confusion Matrix