```
1 import os
 2 import pandas as pd
 3 import numpy as np
 4 import librosa
 5 import random
 6 import time
 7 import pickle
 9 from librosa.display import specshow, waveplot
10
11 from sklearn.preprocessing import MinMaxScaler
12 from sklearn.model selection import train test split
13 from sklearn.preprocessing import LabelBinarizer
14
15 import IPython.display as ipd
16
17 import matplotlib
18 import matplotlib.pyplot as plt
19
20 np.random.seed(42)
21
22 %matplotlib inline
 1 from keras.models import Sequential
 2 from keras.layers import Dense, MaxPooling2D, Conv2D, Flatten, Dropout, Input, BatchNori
 3 from keras.models import Model, load model
 4 from keras.callbacks import Callback, EarlyStopping
 5 from keras.metrics import top k categorical accuracy
   Using TensorFlow backend.
    The default version of TensorFlow in Colab will soon switch to TensorFlow 2.x.
    We recommend you upgrade now or ensure your notebook will continue to use TensorFlow 1.x via the
    %tensorflow_version 1.x magic: more info.
 1 from google.colab import drive
 2 drive.mount('/gdrive')
                                  otebooks
 Saved successfully!
                                   https://accounts.google.com/o/oauth2/auth?client_id=9473189
    Enter your authorization code:
    Mounted at /gdrive
    /gdrive/My Drive/Colab Notebooks
```

Making Sense of Genres

The first step is seeing how many tracks per genre we have in our dataset, and simplifying the output of our neural net to include only the top 5.

Previously, I attempted outputing its confidence for all 161 total genres, and only reached 20% accuracy. You can see that journey in my other notebook, training2_svc.

```
1 genres = pd.read_csv("genres.csv", index_col=0)
2 genres
```

	#tracks	parent	title	top_level
genre_id				
1	8693	38	Avant-Garde	38
2	5271	0	International	2
3	1752	0	Blues	3
4	4126	0	Jazz	4
5	4106	0	Classical	5
•••				
1032	60	102	Turkish	2
1060	30	46	Tango	2
1156	26	130	Fado	2
1193	72	763	Christmas	38
1235	14938	0	Instrumental	1235

163 rows × 4 columns

₽

```
1 genres = genres.sort_values(by='#tracks', ascending=False)
2 genres.head(5)
```

	₽		#tracks	parent	title	top_level
		genre_id				
	Savos	l successfu	illyt	×	xperimental	38
L	Saved successfully!		лпу:		Electronic	15
		12	32923	0	Rock	12
		1235	14938	0	Instrumental	1235
		10	13845	0	Рор	10

Adding Echonest Attributes

Whoop, our top genres are: Experimental, Electronic, Rock, Instrumental, and Pop.

Next, since I want to use this classifier for my senior design project as well, I want to incorporate attributes from *echonest*.

Echnoest, now Spotify, includes numerical values for tracks for traits like dancebility, energy, speechiness, etc-- these will be very valuable when teaching a stick figure to dance. (my senior design)

```
1 echonest = pd.read_csv("echonest.csv", header=[0, 2], skipinitialspace=True, index_col=
2 echonest.head()
```

₽		echonest						
		acousticness	danceability	energy	instrumentalness	liveness	speechiness	t
	track_id							
	2	0.416675	0.675894	0.634476	0.010628	0.177647	0.159310	1
	3	0.374408	0.528643	0.817461	0.001851	0.105880	0.461818	1
	5	0.043567	0.745566	0.701470	0.000697	0.373143	0.124595	1
	10	0.951670	0.658179	0.924525	0.965427	0.115474	0.032985	1

0.019443

0.096567

0.525519 1

5 rows × 249 columns

0.452217

134

 \Box

0.513238 0.560410

	acousticness	danceability	energy	instrumentalness	liveness	speechiness	
track_id							
2	0.416675	0.675894	0.634476	0.010628	0.177647	0.159310	1
3	0.374408	0.528643	0.817461	0.001851	0.105880	0.461818	1
5	0.043567	0.745566	0.701470	0.000697	0.373143	0.124595	1
10	0.951670	0.658179	0.924525	0.965427	0.115474	0.032985	1
134	0.452217	0.513238	0.560410	0.019443	0.096567	0.525519	1

Adding Track Data

Now let's incorporating part of the track dataset.

```
1 tracks = pd.read_csv("tracks.csv", header=[0, 1], skipinitialspace=True, index_col=0)
2 tracks.columns = tracks.columns.droplevel(0)
```

³ tracks.head()

li	information	id	favorites	engineer	date_released	date_created	comments	[→	
								track_id	
		1	4	NaN	2009-01-05 00:00:00	2008-11-26 01:44:45	0	2	
		1	4	NaN	2009-01-05 00:00:00	2008-11-26 01:44:45	0	3	
		1	4	NaN	2009-01-05 00:00:00	2008-11-26 44:45	ılly!	Saved successful	
	NaN	6	4	NaN	2008-02-06 00:00:00	2008-11-26 01:45:08	0	10	
	"spiritual songs" from Nicky Cook	4	2	NaN	2009-01-06 00:00:00	2008-11-26 01:45:05	0	20	

¹ tracks_sub = tracks[['listens', 'name', 'duration', 'genre_top', 'genres', 'title']]

² tracks_sub.head()

	listens	listens	name	duration	<pre>genre_top</pre>	genres	title	title
track_id								
2	6073	1293	AWOL	168	Нір-Нор	[21]	AWOL - A Way Of Life	Food
3	6073	514	AWOL	237	Нір-Нор	[21]	AWOL - A Way Of Life	Electric Ave
5	6073	1151	AWOL	206	Hip-Hop	[21]	AWOL - A Way Of Life	This World
10	47632	50135	Kurt Vile	161	Pop	[10]	Constant Hitmaker	Freeway
20	2710	361	Nicky Cook	311	NaN	[76, 103]	Niris	Spiritual Level

¹ tracks_sub.columns = ['listens_album', 'listens_track', 'name', 'duration', 'genre_top'

₽

₽		listens_album	listens_track	name	duration	genre_top	genres	title_album	1
	track_id								
	2	6073	1293	AWOL	168	Нір-Нор	[21]	AWOL - A Way Of Life	
	3	6073	514	AWOL	237	Hip-Hop	[21]	AWOL - A Way Of Life	
	5	6073	1151	AWOL	206	Hip-Hop	[21]	AWOL - A Way Of Life	
	10	47632	50135	Kurt Vile	161	Pop	[10]	Constant Hitmaker	
	20	2710	361	Nicky Cook	311	NaN	[76, 103]	Niris	

Saved successfully!

Merging Tracks, Echonest, and Genres

oh boy

```
1 tracks_echo = pd.merge(tracks_sub, echonest_sub, how="inner", on="track_id")
```

₽

¹ tracks_sub.head()

¹ tracks_echo.head()

	_	_		•		-	_
track_id							
2	6073	1293	AWOL	168	Нір-Нор	[21]	AWOL - A Way Of Life
3	6073	514	AWOL	237	Нір-Нор	[21]	AWOL - A Way Of Life
5	6073	1151	AWOL	206	Нір-Нор	[21]	AWOL - A Way Of Life
10	47632	50135	Kurt Vile	161	Pop	[10]	Constant Hitmaker
134	6073	943	AWOL	207	Нір-Нор	[21]	AWOL - A Way Of Life

listens album listens track name duration genre top genres title album 1

¹ tracks_echo_genres.head()

Г→	listens_album	listens_track	name	duration	genre_top	genres	title_album	1
track_id								
2	6073	1293	AWOL	168	Нір-Нор	[21]	AWOL - A Way Of Life	
3	6073	514	AWOL	237	Hip-Hop	[21]	AWOL - A Way Of Life	
5	6073	1151	AWOL	206	Hip-Hop	[21]	AWOL - A Way Of Life	
10	47632	50135	Kurt Vile	161	Pop	[10]	Constant Hitmaker	
124 Saved successful	6073 y	943 ×	AWOL	207	Нір-Нор	[21]	AWOL - A Way Of Life	

¹ tracks_echo_genres.to_pickle("./tracks_echo_genres.pkl")

Adding Features

This is the final piece left to merge into our monster dataset. There are a lot of attributes here-- 518-- so I want to do some dimensionality reduction here. I will be using PCA post-merge.

¹ tracks_echo_genres = pd.merge(tracks_echo, genres, how="left", left_on="genre_top", rigi

¹ features = pd.read_csv("features.csv", header=[0, 1, 2], skipinitialspace=True, index_cc

² features.head()

₽	feature	chroma_c	a_cens							
	statistics	kurtosis								
	number	01	02	03	04	05	06	07	08	09
	track_id									
	2	7.180653	5.230309	0.249321	1.347620	1.482478	0.531371	1.481593	2.691455	0.86
	3	1.888963	0.760539	0.345297	2.295201	1.654031	0.067592	1.366848	1.054094	0.10
	5	0.527563	-0.077654	-0.279610	0.685883	1.937570	0.880839	-0.923192	-0.927232	0.66
	10	3.702245	-0.291193	2.196742	-0.234449	1.367364	0.998411	1.770694	1.604566	0.52
	20	-0.193837	-0.198527	0.201546	0.258556	0.775204	0.084794	-0.289294	-0.816410	0.04

5 rows × 518 columns

/usr/local/lib/python3.6/dist-packages/pandas/core/reshape/merge.py:617: UserWarning: me warnings.warn(msg, UserWarning)

1 monster.head()

 \Box

		listens_album	listens_track	name	duration	genre_top	genres	title_album	1
	track_id								
	2	6073	1293	AWOL	168	Hip-Hop	[21]	AWOL - A Way Of Life	
	3	6073	514	AWOL	237	Нір-Нор	[21]	AWOL - A Way Of Life	
L	Saved successful	ly!	× 1151	AWOL	206	Hip-Hop	[21]	AWOL - A Way Of Life	
	10	47632	50135	Kurt Vile	161	Рор	[10]	Constant Hitmaker	
	134	6073	943	AWOL	207	Нір-Нор	[21]	AWOL - A Way Of Life	

5 rows × 539 columns

^{1 #} MERGING!!!

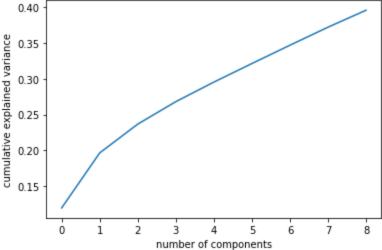
² monster = pd.merge(tracks_echo_genres, features, how="inner", on="track_id")

¹ monster.to_pickle("./monster.pkl")

→ PCA Shenaniaans for Dimensionality Reduction

```
1 from sklearn.preprocessing import StandardScaler
  2 feats = monster.columns
  4 # Separating out the features
  5 numerics = ['int16', 'int32', 'int64', 'float16', 'float32', 'float64']
  6 x = monster.select dtypes(include=numerics).values
  8 # x = monster.loc[:, feats].values
  9 # Separating out the target
10 y = monster.loc[:,['genre_top']].values
12 # Standardizing the features
13 X = StandardScaler().fit transform(x)
 1 X.shape
          (13129, 532)
 \Gamma
  1 # no nan vals allowed!!!
  2 from sklearn.impute import SimpleImputer
  4 imputer = SimpleImputer(missing_values=np.nan, strategy='mean')
  5 imputer = imputer.fit(X[:,1:532])
  6 \times [:, 1:532] = imputer.transform(X[:, 1:532])
  1 from sklearn.decomposition import PCA
  3 pca = PCA(n_components=9) # for number of big attributes?
  4 principalComponents = pca.fit transform(X)
  5 principalDf = pd.DataFrame(data = principalComponents
                                           , columns = ['principal component 1', 'principal component 2', 'principal component 2', 'principal component 1', 'principal component 2', 'principal component 1', 'princip
  1 # SCREE PLOT
  2 print(pca.explained_variance_ratio_)
  3 print(np.cumsum(pca.explained variance ratio ))
  Saved successfully!
 opic.pioc(np.cumbum(pca.cxpiuined_variance_ratio_))
  7 plt.xlabel('number of components')
  8 plt.ylabel('cumulative explained variance')
  9 plt.show()
 \Box
```

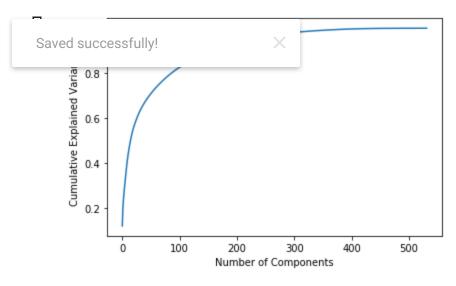
```
[0.11959576 0.07689131 0.04026169 0.03136196 0.02734066 0.0259367 0.02560827 0.02520922 0.02367956]
[0.11959576 0.19648707 0.23674876 0.26811072 0.29545139 0.32138809 0.34699636 0.37220558 0.39588514]
```



▼ Ok!!!!!!!! So, this shows that with 9 components, it represents 40% of variance in the data.

Let's make a generic scree plot to see how many components might make more sense:

```
1 # SCREE PLOT
2
3 #Explained variance
4 pca = PCA().fit(X)
5 plt.plot(np.cumsum(pca.explained_variance_ratio_))
6 plt.xlabel('Number of Components')
7 plt.ylabel('Cumulative Explained Variance')
8
9 plt.plot(250, np.cumsum(pca.explained_variance_ratio_)[250], "ob")
10 plt.plot(200, np.cumsum(pca.explained_variance_ratio_)[200], "or")
11 plt.plot(150, np.cumsum(pca.explained_variance_ratio_)[150], "og")
12
13 plt.show()
```



This plot shows us that around 150-250 components might be a better number to try. Let's try it:

```
1 #Explained variance
 2
 3 fig, axs = plt.subplots(3)
 4 fig.suptitle('Comparing Num of PCA Components')
 5 # fig.xlabel('Number of Components')
 6 # fig.ylabel('Cumulative Explained Variance')
 8 # PLOT 1: 150 COMPONENTS
 9 pca = PCA(n components=150)
10 principalComponents = pca.fit_transform(X)
11 col names = [("col " + str(i)) for i in range(150)]
12 principalDf = pd.DataFrame(data = principalComponents
                , columns = col_names)
14 axs[0].plot(np.cumsum(pca.explained variance ratio ))
15
16 # PLOT 2: 200 COMPONENTS
17 pca = PCA(n components=200)
18 principalComponents = pca.fit_transform(X)
19 col names = [("col " + str(i)) for i in range(200)]
20 principalDf = pd.DataFrame(data = principalComponents
                , columns = col names)
22 axs[1].plot(np.cumsum(pca.explained_variance_ratio_))
24 # PLOT 3: 250 COMPONENTS
25 pca = PCA(n components=250)
26 principalComponents = pca.fit transform(X)
27 col_names = [("col_" + str(i)) for i in range(250)]
28 principalDf = pd.DataFrame(data = principalComponents
                , columns = col names)
30 axs[2].plot(np.cumsum(pca.explained_variance_ratio_))
    [<matplotlib.lines.Line2D at 0x7f5f3d5f8978>]
               Comparing Num of PCA Components
     0.75
     0.50
     0.25
                                   100
 Saved successfully!
                             100
      1.0
      0.5
                  50
                         100
                                150
                                        200
                                                250
```

So.... they are looking basically the same! Let's see if we can go smaller than 150 components so that we can be faster while training

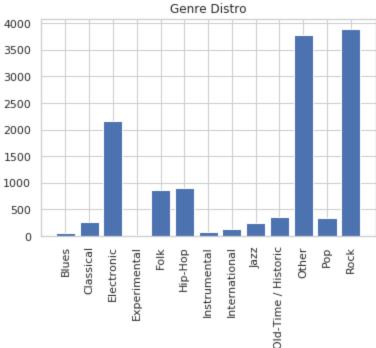
```
1 #Explained variance
2 pca = PCA(n_components=100) # for number of big attributes?
3 principalComponents = pca.fit transform(X)
```

```
4 col_names = [("col_" + str(i)) for i in range(100)]
 5 principalDf = pd.DataFrame(data = principalComponents
                 , columns = col_names)
 7
 8 plt.plot(np.cumsum(pca.explained_variance_ratio_))
 9 plt.xlabel('Number of Components')
10 plt.ylabel('Cumulative Explained Variance')
11 plt.show()
13 print(100, np.cumsum(pca.explained variance ratio )[99])
\Box
       0.8
     Cumulative Explained Variance
       0.7
       0.6
       0.5
       0.4
       0.3
       0.2
       0.1
                   20
                            40
                                    60
                                                   100
                        Number of Components
    100 0.8235167577314023
 1 def compare_n_comp(n):
 2
    pca = PCA(n_components=n)
 3
     principalComponents = pca.fit transform(X)
 4
     print(n, str((np.cumsum(pca.explained_variance_ratio_)[n-1])*100), str('%'))
 5
 6 compare_n_comp(150)
 7 compare_n_comp(175)
 8 compare n comp(200)
    150 89.51656816113069 %
    175 91.96887994360743 %
    200 93.89739765751429 %
 Saved successfully!
                                    ts. That'll be a 62% reduction!
 1 #Explained variance
 2 pca = PCA(n components=200)
 3 principalComponents = pca.fit_transform(X)
 4 col_names = [("col_" + str(i)) for i in range(200)]
 5 principalDf = pd.DataFrame(data = principalComponents
 6
                 , columns = col_names)
 1 print(principalDf.shape)
 2 print(y.shape)
```

I want to visualize the data I'm going to train with in some way... let's try taking the mean of all the numerical values in our PCA components and map them to their respective genre.

```
1 y_df = pd.DataFrame(y)
2 y_df = y_df.replace(np.nan, 'Other', regex=True)
3 y_map = dict(zip(*np.unique(y_df, return_counts=True)))
4 plt.bar(y_map.keys(), y_map.values())
5 plt.xticks(rotation='vertical')
6 plt.title("Genre Distro")
7
8 print(y_map)
```

[→ {'Blues': 66, 'Classical': 265, 'Electronic': 2170, 'Experimental': 17, 'Folk': 874, 'Hi



Time to Train!

```
8 data_test = scaler.transform(data_test)

1 pca = PCA(.93) # this is about 200 components as we saw earlier
2 pca.fit(data_train)

C> PCA(copy=True, iterated_power='auto', n_components=0.93, random_state=None, svd solver='auto', tol=0.0, whiten=False)
```

Bad

 \Box

```
→ 4 cells hidden
```

So there's a problem here: I'm trying to use string values as labels for my dataset, which is not allowed. I decided to try this route (without thinking it all the way through) because the numerical genre values, genre_ids, were stored as a string of a list of a list, and I wanted to try to avoid dealing with that mess. Looks like it's unavoidable so let's...

Deal with the genre Label Mess

```
1 with open("genre_labels.pkl", "rb") as handle:
2   genre_labels = pickle.load(handle)
3
4 genrelabels = pd.DataFrame.from_dict(genre_labels)
5 genrelabels.head()
```

С→		genre_id	<pre>genre_title</pre>
	0	1	Avant-Garde
	1	2	International
	2	3	Blues
	3	4	Jazz

```
Saved successfully!
```

```
1 abel_train2 = pd.merge(label_train, genrelabels, how="left", left_on=0, right_on="genre")
1 label_train2.head()
```

```
0 genre_id genre_title
    0 Other
                  NaN
                                NaN
      Other
                  NaN
                                NaN
    1
    2
       Rock
                  12.0
                                Rock
    3
       Rock
                  12.0
                                Rock
      Other
                  NaN
                                NaN
1 abel_test2 = pd.merge(label_test, genrelabels, how="left", left_on=0, right_on="genre_ti
2 abel_test2.head()
          0 genre_id genre_title
    0 Other
                  NaN
                                NaN
       Other
                  NaN
                                NaN
    2
       Rock
                  12.0
                                Rock
    3
      Other
                  NaN
                                NaN
    4 Other
                  NaN
                                NaN
1 label_test = label_test2["genre_id"]
2 label_train = label_train2["genre_id"]
3 label test = label test.replace(np.nan, 0, regex=True)
4 label_train = label_train.replace(np.nan, 0, regex=True)
6 label_test, label_train
   (0
              0.0
              0.0
    1
    2
             12.0
    3
              0.0
    4
              0.0
Saved successfully!
    4330
             12.U
    4331
             12.0
              4.0
    Name: genre id, Length: 4333, dtype: float64, 0
                                                             12.0
            17.0
    2
            12.0
    3
            17.0
            17.0
            . . .
    196
            12.0
    197
            12.0
    198
            12.0
    199
            21.0
```

 \Box

5

 \Box

200

12.0

Name: genre_id, Length: 201, dtype: float64)

→ OK! let's try to train again..

```
1 one_hot_train_labels = to_categorical(label_train.values)
 2 one hot test labels = to categorical(label test.values)
 4 one hot train labels.shape, one hot train labels.shape
    ((201, 22), (201, 22))
1 from keras.models import Sequential
2 from keras.layers import Dense, Activation
 4 model = Sequential()
5 model.add(Dense(32, activation='relu', input shape=(200, )))
 6 model.add(Dense(64, activation='relu'))
 7 model.add(Dense(64, activation='relu'))
8 model.add(Dense(32, activation='tanh'))
10 model.add(Dense(161, activation='softmax'))
11
12 \text{ output} = 164
13 model.add(Dense(output, activation='sigmoid')) # all genres
14
15 model.compile(optimizer='rmsprop', loss='sparse categorical crossentropy', metrics=['acc
16 model.summary()
    Model: "sequential 53"
                                                           Param #
    Layer (type)
                                 Output Shape
    ______
    dense 253 (Dense)
                                                           6432
                                 (None, 32)
                                                           2112
    dense 254 (Dense)
                                 (None, 64)
    dense 255 (Dense)
                                 (None, 64)
                                                           4160
    dense 256 (Dense)
                                 (None, 32)
                                                           2080
 Saved successfully!
                                 (None, 161)
                                                           5313
    dense 258 (Dense)
                                 (None, 164)
                                                           26568
    Total params: 46,665
    Trainable params: 46,665
    Non-trainable params: 0
 1 model.fit(data train, label train, epochs=35, batch size=512, validation split=0.1)
```

С→

```
Epoch 7/35
180/180 [===============] - 0s 51us/step - loss: 4.5861 - acc: 0.7611 - v
Epoch 8/35
Epoch 9/35
Epoch 10/35
Epoch 11/35
Epoch 12/35
Epoch 13/35
Epoch 14/35
Epoch 15/35
Epoch 16/35
Epoch 17/35
Epoch 18/35
Epoch 19/35
Epoch 20/35
Epoch 21/35
Epoch 22/35
Epoch 23/35
Epoch 24/35
Epoch 25/35
Epoch 26/35
Epoch 27/35
Epoch 28/35
      =======] - 0s 60us/step - loss: 4.5468 - acc: 0.7611 - v
Saved successfully!
      =======] - 0s 70us/step - loss: 4.5449 - acc: 0.7611 - v
Epoch 30/35
Epoch 31/35
Epoch 32/35
Epoch 33/35
Epoch 34/35
Epoch 35/35
```

```
1 # serialize model to JSON
2 model_json = model.to_json()
3 with open("model_1.json", "w") as json_file:
4     json_file.write(model_json)
5 # serialize weights to HDF5
6 model.save_weights("model_1.h5")
7 print("Saved model to disk")

□→ Saved model to disk
```

Not bad, we got to 76% (model_1) which is much better than the 21% we were getting before! Let's play with a different regularizer to see if we can do better..

```
1 com keras.models import Sequential
 2 com keras.layers import Dense, Activation
3 com keras import optimizers
5 >del = Sequential()
 6 >del.add(Dense(32, activation='relu', input shape=(200, )))
 7 >del.add(Dense(64, activation='relu'))
8 >del.add(Dense(128, activation='relu'))
9 model.add(Dropout(0.1))
10 model.add(Dense(128, activation='relu'))
11 >del.add(Dense(64, activation='relu'))
12 odel.add(Dense(32, activation='relu'))
13 >del.add(Dense(161, activation='relu'))
 Saved successfully!
                                 ion='sigmoid')) # all genres
17
18 Jd = optimizers.SGD(lr=1e-08, clipvalue=1)
19 >del.compile(optimizer=sgd, loss='sparse categorical crossentropy', metrics=['accuracy']
20 odel.summary()
\Box
```

Layer (typ	e)	Output	Shape	Param #
dense_168	(Dense)	(None,	32)	6432
dense_169	(Dense)	(None,	64)	2112
dense_170	(Dense)	(None,	128)	8320
dense_171	(Dense)	(None,	64)	8256
donco 172	(Dongo)	/Mono	331	2000

1 model.fit(data train, label train, epochs=20, batch size=512, validation split=0.33)

```
Train on 134 samples, validate on 67 samples
Epoch 1/20
Epoch 2/20
Epoch 3/20
Epoch 4/20
Epoch 5/20
Epoch 6/20
Epoch 7/20
Epoch 8/20
Epoch 9/20
Epoch 10/20
Epoch 11/20
Epoch 12/20
Epoch 13/20
Epoch 14/20
      =======] - 0s 83us/step - loss: 5.1049 - acc: 0.0075 - v
Saved successfully!
     ----========] - 0s 68us/step - loss: 5.1049 - acc: 0.0075 - v
134/134 [---
Epoch 16/20
Epoch 17/20
Epoch 18/20
Epoch 19/20
Epoch 20/20
```

1	
Saved successfully!	×

Nope. Let's just stick with the weights from before.