## IMDB\_RNN

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```
library(keras)
library(reticulate)
library(dplyr)

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##
## filter, lag

## The following objects are masked from 'package:base':
##
## intersect, setdiff, setequal, union
```

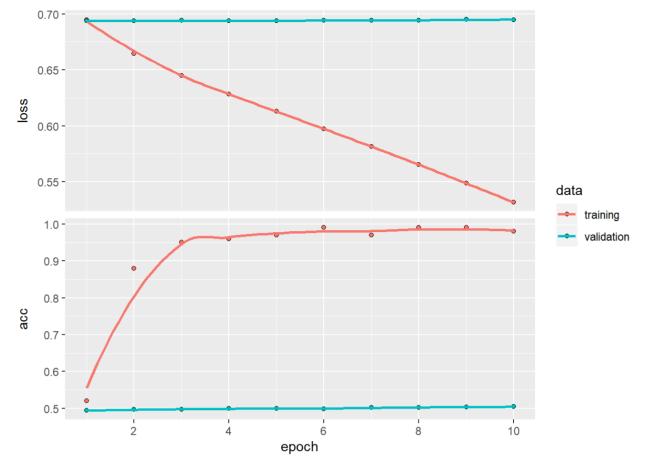
```
imdb_dir <- "D:/MSBA/Adv ML/aclImdb"</pre>
train_dir <- file.path(imdb_dir, "train")</pre>
labels <- c()
texts <- c()
for (label_type in c("neg", "pos")) {
  label <- switch(label type, neg = 0, pos = 1)</pre>
  dir_name <- file.path(train_dir, label_type)</pre>
 for (fname in list.files(dir_name, pattern = glob2rx("*.txt"),
                            full.names = TRUE)) {
    texts <- c(texts, readChar(fname, file.info(fname)$size))</pre>
    labels <- c(labels, label)</pre>
}
maxlen <- 150
                               # We will cut reviews after 150 words
training samples <- 100  # We will be training on 200 samples
validation_samples <- 10000  # We will be validating on 10000 samples
max_words <- 10000
                              # We will only consider the top 10,000 words in the dataset
# tokenizing the words
tokenizer <- text_tokenizer(num_words = max_words) %>%
 fit text tokenizer(texts)
sequences <- texts_to_sequences(tokenizer, texts)</pre>
word_index = tokenizer$word_index
# Turns the list of integers into a 2D integer tensor shape (samples,maxlen)
data <- pad sequences(sequences, maxlen = maxlen)</pre>
labels <- as.array(labels)</pre>
cat("Shape of data tensor:", dim(data), "\n")
```

```
## Shape of data tensor: 25000 150
```

```
#Shape of data tensor: 25000 150 cat('Shape of label tensor:', dim(labels), "\n")
```

## Shape of label tensor: 25000

```
#Shape of label tensor: 25000
set.seed(123)
indices <- sample(1:nrow(data))</pre>
training indices <- indices[1:training samples]</pre>
validation_indices <- indices[(training_samples + 1):</pre>
                                  (training samples + validation samples)]
train_data <- data[training_indices,]</pre>
train_label <- labels[training_indices]</pre>
valid_data <- data[validation_indices,]</pre>
valid_label<- labels[validation_indices]</pre>
test dir <- file.path(imdb dir, "test")</pre>
labels <- c()
texts <- c()
for (label_type in c("neg", "pos")) {
  label <- switch(label_type, neg = 0, pos = 1)</pre>
  dir_name <- file.path(test_dir, label_type)</pre>
  for (fname in list.files(dir_name, pattern = glob2rx("*.txt"),
                             full.names = TRUE)) {
    texts <- c(texts, readChar(fname, file.info(fname)$size))</pre>
    labels <- c(labels, label)</pre>
  }
}
sequences <- texts_to_sequences(tokenizer, texts)</pre>
x_test <- pad_sequences(sequences, maxlen = maxlen)</pre>
y_test <- as.array(labels)</pre>
# Using an embedding layer and classifier on the IMDB data
model <- keras model sequential() %>% layer embedding(input dim = 10000,output dim = 8,input length =
maxlen) %>%
  layer_flatten() %>% layer_dense(units=1,activation = "sigmoid")
model %>% compile(optimizer = "rmsprop",loss = "binary_crossentropy",metrics=c("acc"))
history <- model %>% fit(train_data,train_label,epochs=10,batch_size=32,validation_data = list(valid_
data,valid_label))
# Plot of Accuracy and Loss function of the model
plot(history)
```



# From the above plot we can see that the model is performing with ~50% accuracy with 100 samples in the training data While the Validation dataset has 1000 samples.

```
# validation accuracy

# Evaluating the test dataset
model %>% fit(
    train_data,
    train_label,
    epochs = 2,
    batch_size = 32)
result <- model %>% evaluate(x_test,y_test)
result
```

```
## $loss
## [1] 0.69567
##
## $acc
## [1] 0.5
```

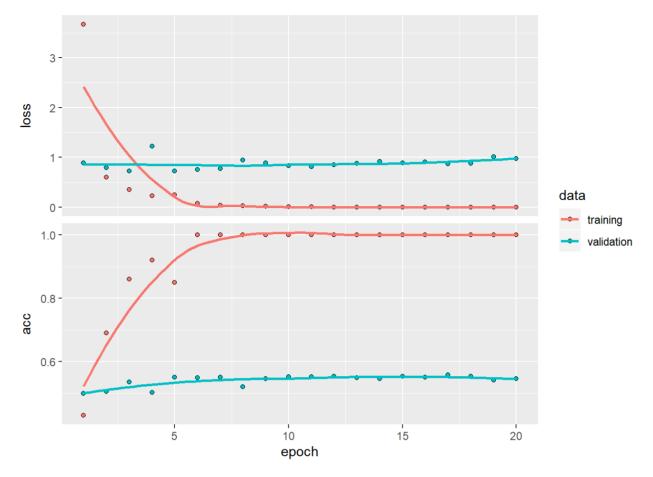
```
#The Test accuracy of the model is 0.50

# Parsing the GLoVe word-embeddings file
glove_dir = 'D:/MSBA/Adv ML/glove.6B'
lines <- readLines(file.path(glove_dir, "glove.6B.100d.txt"))

embeddings_index <- new.env(hash = TRUE, parent = emptyenv())
for (i in 1:length(lines)) {
    line <- lines[[i]]
    values <- strsplit(line, " ")[[1]]
    word <- values[[1]]
    embeddings_index[[word]] <- as.double(values[-1])
}
cat("Found", length(embeddings_index), "word vectors.\n")</pre>
```

## Found 400000 word vectors.

```
# Preparing the GloVe word-embeddings matrix
embedding dim <- 100
embedding_matrix <- array(0, c(max_words, embedding_dim))</pre>
for (word in names(word_index)) {
  index <- word_index[[word]]</pre>
  if (index < max words) {</pre>
    embedding_vector <- embeddings_index[[word]]</pre>
    if (!is.null(embedding vector))
      embedding matrix[index+1,] <- embedding vector
}
# Model construction
model <- keras_model_sequential() %>%
  layer embedding(input dim = max words, output dim = embedding dim,input length = maxlen) %>%
  layer flatten() %>%
  layer_dense(units = 32, activation = "relu") %>%
  layer dense(units = 1, activation = "sigmoid")
# Loading pretrained word embeddings into the embedding layer
get_layer(model, index = 1) %>%
  set_weights(list(embedding_matrix)) %>%
  freeze weights()
model %>% compile(
  optimizer = "rmsprop",
  loss = "binary_crossentropy",
  metrics = c("acc")
)
history1 <- model %>% fit(
 train_data, train_label,
  epochs = 20,
  batch_size = 32,
  validation_data = list(valid_data , valid_label)
plot(history1)
```



# Trying to improve the model performance using pretrained networks but we observe that model still performs with ~50% accuracy that is due to less number of samples in training data. As it does not see more patterns in the data.

```
# validation accuracy is 55.81 by cutting reviews for first 150 words in every 100 samples

model %>% fit(
   train_data, train_label,
   epochs = 2,
   batch_size = 32)

result1 <- model %>% evaluate(x_test,y_test)
   result1 # Test Accuracy of the model is 55%
```

```
## $loss
## [1] 0.9918165
##
## $acc
## [1] 0.54552
```