▼ IMPORTING LIBRARY

```
import numpy as np
import pandas as pd
import re
import os
import json
import pickle
from tqdm import tqdm
import glob
import matplotlib.pyplot as plt
from bs4 import BeautifulSoup
from keras.preprocessing.text import Tokenizer
from keras.utils import pad_sequences
from keras_preprocessing.sequence import pad_sequences
from nltk.corpus import stopwords
from tensorflow.keras.layers import Input, LSTM, Embedding, Dense, Concatenate, TimeDistribut
from tensorflow.keras.models import Model
from tensorflow.keras.callbacks import EarlyStopping
import warnings
pd.set option("display.max colwidth", 200)
warnings.filterwarnings("ignore")
from google.colab import drive
drive.mount('/content/drive')
     Mounted at /content/drive
```

▼ TAKING SEGMENTED DATASET

```
Editorial.
3
Editorial.
Editorial.
. . .
. . .
9995
                                  Video-Observed Therapy With a Notification System
for Improving the Monitoring of Tuberculosis Treatment in Thailand: Usability Study
9996
         Modelling the impact of the coronavirus pandemic on bowel cancer screening
outcomes in England: A decision analysis to prepare for future screening disruption
      Epidemiology, Clinical Features, and Outcomes of Multisystem Inflammatory
Syndrome in Children (MIS-C) and Adolescents—a Live Systematic Review and Meta-
analysis
9998
The Molecular Basis of Resilience: A Narrative Review
                                                                         "Exams by
You": Having Students Write and Complete Their Own Exams During the COVID-19
Pandemic
abstract
0
[This corrects the article on p. 147 in vol. 33.].
NaN
2
NaN
3
NaN
4
NaN
. . .
. . .
9995 BACKGROUND: In Thailand, the health care system has struggled to cope with
COVID-19, resulting in directly observed therapy for tuberculosis being de-
emphasized. A video-observed therapy (VOT) sys...
9996 The English Bowel Cancer Screening Programme invites people between the ages
of 60 and 74 to take a Faecal Immunochemical Test every two years. This programme
was interrupted during the coronaviru...
9997 PURPOSE OF REVIEW: A multisystem inflammatory condition occurring in children
and adolescents with COVID-19 has become increasingly recognized and widely studied
globally. This review aims to inve...
9998 Resilience refers to the adaptability of a person - an ability to "bounce-
back" from stressors. We question if resilience can be strengthened, potentially to
decrease the risk of stress-related di...
9999 BACKGROUND: The COVID-19 pandemic made it difficult to proctor exams after the
forced transition to remote teaching and learning. OBJECTIVE: We evaluated students'
experiences creating and answeri...
```

df.head()

data.

	title	abstract	1
0	Correction.	[This corrects the article on p. 147 in vol. 33.].	
1	Correspondence.	NaN	
2	Editorial.	NaN	
3	Editorial.	NaN	

→ DATA PREPROCESSING

```
df.isna().sum()
    title     1
    abstract    1184
    dtype: int64

df.drop_duplicates(subset=['abstract'],inplace=True)#dropping duplicates
df.dropna(axis=0,inplace=True)#dropping na

data = df[['title','abstract']]

data.head()
```

	title	abstract
0	Correction.	[This corrects the article on p. 147 in vol. 33.].
11	A bioactive compliant vascular graft modulates macrophage polarization and maintains patency with robust vascular remodeling.	Conventional synthetic vascular grafts are associated with significant failure rates due to their mismatched mechanical properties with the native vessel and poor regenerative potential. Though di
12	Strategies to increase rural maternal utilization of skilled health personnel for childbirth delivery in low- and middle-income countries: a narrative review.	BACKGROUND Skilled attendance at birth is considered key to accomplishing Sustainable Development Goal (SDG) 3.1 aimed at reducing maternal mortality. Many maternal deaths can be prevented if a wo
13	A 'Multiomic' Approach of Saliva Metabolomics, Microbiota, and Serum Biomarkers to Assess the Need of Hospitalization in Coronavirus Disease	Background and Aims The SARS-CoV-2 pandemic has overwhelmed the treatment capacity of the health care systems during the highest viral diffusion rate. Patients reaching the emergency department ha
info	()	
Int6	ss 'pandas.core.frame.DataFrame'> 4Index: 8810 entries, 0 to 9999 columns (total 2 columns): Column Non-Null Count Dtype	

```
0
          title
                    8810 non-null
                                    object
          abstract 8810 non-null
      1
                                    object
     dtypes: object(2)
     memory usage: 206.5+ KB
from wordcloud import WordCloud, STOPWORDS
# Thanks : https://www.kaggle.com/aashita/word-clouds-of-various-shapes ##
def plot wordcloud(text, mask=None, max words=200, max font size=50, figure size=(15.0,15.0),
                   title = None, title size=20, image color=False,color = None):
   stopwords = set(STOPWORDS)
   more stopwords = {'one', 'br', 'Po', 'th', 'sayi', 'fo', 'Unknown'}
   stopwords = stopwords.union(more stopwords)
   wordcloud = WordCloud(background color=color,
                    stopwords = stopwords,
                    max words = max words,
                    max font size = max font size,
                    random state = 42,
                    width=800,
                    height=400,
                    mask = mask)
   wordcloud.generate(str(text))
   plt.figure(figsize=figure_size)
   if image color:
        image_colors = ImageColorGenerator(mask);
        plt.imshow(wordcloud.recolor(color_func=image_colors), interpolation="bilinear");
        plt.title(title, fontdict={'size': title size,
                                  'verticalalignment': 'bottom'})
   else:
        plt.imshow(wordcloud);
        plt.title(title, fontdict={'size': title size, 'color': 'black',
                                  'verticalalignment': 'bottom'})
   plt.axis('off');
   plt.tight layout()
plot wordcloud(data['title'].values, title="Word Cloud of Title",color = 'black')
plot wordcloud(data['abstract'].values, title="Word Cloud of Abstarct",color = 'black')
```

Word Cloud of Title

```
middlegraft

Basis Vascular Children

Exams

Strategies Epidemiology health

Strategies Epidemiology health

Strategies Epidemiology health

Strategies Epidemiology health

Income increase increase increase increase increase complete Syndrome increase inc
```

Word Cloud of Abstarct



▼ DATA CLEANING

```
# A list of contractions from http://stackoverflow.com/questions/19790188/expanding-english-l
contraction mapping = {"ain't": "is not", "aren't": "are not", "can't": "cannot", "'cause": "b
                           "didn't": "did not", "doesn't": "does not", "don't": "do not", "h
                           "he'd": "he would", "he'll": "he will", "he's": "he is", "how'd": "
                           "I'd": "I would", "I'd've": "I would have", "I'll": "I will", "I'l
                           "i'd've": "i would have", "i'll": "i will", "i'll've": "i will ha
                           "it'd've": "it would have", "it'll": "it will", "it'll've": "it wi
                           "mayn't": "may not", "might've": "might have", "mightn't": "might n
                           "mustn't": "must not", "mustn't've": "must not have", "needn't": "
                           "oughtn't": "ought not", "oughtn't've": "ought not have", "shan't"
                           "she'd": "she would", "she'd've": "she would have", "she'll": "she
                           "should've": "should have", "shouldn't": "should not", "shouldn't'
                           "this's": "this is", "that'd": "that would", "that'd've": "that wou
                           "there'd've": "there would have", "there's": "there is", "here's":
                           "they'll": "they will", "they'll've": "they will have", "they're":
                           "wasn't": "was not", "we'd": "we would", "we'd've": "we would have
                           "we've": "we have", "weren't": "were not", "what'll": "what will",
                           "what's": "what is", "what've": "what have", "when's": "when is",
                           "where've": "where have", "who'll": "who will", "who'll've": "who
                           "why's": "why is", "why've": "why have", "will've": "will have", "
                           "would've": "would have", "wouldn't": "would not", "wouldn't've":
                           "y'all'd": "you all would","y'all'd've": "you all would have","y'a
                           "you'd": "you would", "you'd've": "you would have", "you'll": "you
                           "you're": "you are", "you've": "you have"}
```

```
import nltk
nltk.download('stopwords')
stop words = set(stopwords.words('english'))
def text cleaner(text,num):
    newString = text.lower()
    newString = BeautifulSoup(newString, "lxml").text
    newString = re.sub(r'\([^{\wedge}]*\)', '', newString)
    newString = re.sub('"','', newString)
    newString = ' '.join([contraction mapping[t] if t in contraction mapping else t for t in
    newString = re.sub(r"'s\b","",newString)
    newString = re.sub("[^a-zA-Z]", " ", newString)
    newString = re.sub('[m]{2,}', 'mm', newString)
    if(num==0):
        tokens = [w for w in newString.split() if not w in stop words]
    else:
        tokens=newString.split()
    long_words=[]
    for i in tokens:
```

```
#removing short word
```

```
long_words.append(i)
return (" ".join(long_words)).strip()

[nltk_data] Downloading package stopwords to /root/nltk_data...
[nltk_data] Unzipping corpora/stopwords.zip.

#call the function
cleaned_text = []
for t in data['abstract']:
    cleaned_text.append(text_cleaner(t,0))
cleaned_text[:3]
```

['corrects article vol',

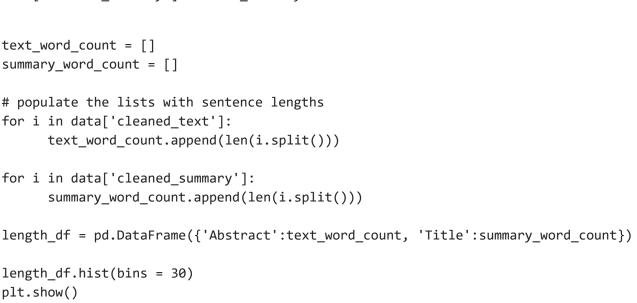
if len(i)>1:

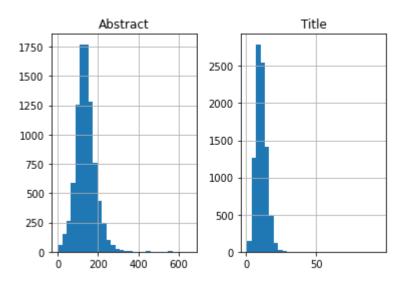
'conventional synthetic vascular grafts associated significant failure rates due mismatched mechanical properties native vessel poor regenerative potential though different tissue engineering approaches used improve biocompatibility synthetic vascular grafts still crucial develop new generation synthetic grafts match dynamics native vessel direct host response achieve robust vascular regeneration size pores within implanted biomaterials shown significant effects macrophage polarization confirmed necessary efficient vascular formation remodeling developed biodegradable autoclavable synthetic vascular grafts new polyurethane elastomer tailored grafts interconnected pore sizes promote macrophage populations pro regenerative phenotype improve vascular regeneration patency rate synthetic vascular grafts showed similar mechanical properties native blood vessels encouraged macrophage populations varying phenotypic expression maintained patency vascular regeneration one month rat carotid interposition model four month rat aortic interposition model innovative bioactive synthetic vascular graft holds promise treat clinical vascular diseases',

'background skilled attendance birth considered key accomplishing sustainable development goal aimed reducing maternal mortality many maternal deaths prevented woman receives care skilled health personnel maternal utilization skilled health delivery services rural areas low middle income countries compared urban areas previous studies found community based interventions may increase rural maternal uptake skilled health delivery services evidence lacking strategies effective objective review effectiveness community based strategies increase rural maternal utilization skilled health personnel childbirth delivery low middle income countries methods conducted narrative review pubmed cinahl cochrane library psycinfo databases searched articles database inception november key search terms pre determined information extracted studies meeting inclusion criteria cluster randomized trials rural setting reproductive aged women community engagement low middle income countries studies considered effective statistically significant narrative synthesis conducted results ten cluster randomized trials candidate citations met inclusion criteria strategies included home based visits women groups combined approaches ten articles three studies found significantly increase maternal uptake skilled health personnel delivery used different strategy results inconclusive strategies effective limitations review include heterogeneity generalizability studies conclusions research suggests different strategies may effective improving maternal utilization skilled health personnel delivery certain rural settings ineffective others research warranted better understand context strategies may effective conditions']

#call the function

```
Incre2 LSTM.ipynb - Colaboratory
cleaned summary = []
for t in data['title']:
    cleaned_summary.append(text_cleaner(t,0))
cleaned summary[:5]
     ['correction',
      'bioactive compliant vascular graft modulates macrophage polarization maintains
     patency robust vascular remodeling',
      'strategies increase rural maternal utilization skilled health personnel childbirth
     delivery low middle income countries narrative review',
      'multiomic approach saliva metabolomics microbiota serum biomarkers assess need
     hospitalization coronavirus disease',
      'chapter infectious disease emergencies']
data['cleaned_text']=cleaned_text
data['cleaned summary']=cleaned summary
text word count = []
summary word count = []
# populate the lists with sentence lengths
for i in data['cleaned text']:
```





data['cleaned summary'] = data['cleaned summary'].apply(lambda x : 'sostok '+ x + ' eostok')

```
max text len = 100
max summary len = 35
from sklearn.model selection import train test split
x_tr,x_val,y_tr,y_val=train_test_split(np.array(data['cleaned_text']),np.array(data['cleaned_
from keras.preprocessing.text import Tokenizer
#prepare a tokenizer for reviews on training data
x tokenizer = Tokenizer()
x tokenizer.fit on texts(list(x tr))
thresh=5
cnt=0
tot cnt=0
freq=0
tot_freq=0
for key,value in x_tokenizer.word_counts.items():
   tot cnt=tot cnt+1
   tot freq=tot freq+value
   if(value<thresh):</pre>
        cnt=cnt+1
        freq=freq+value
print("% of rare words in vocabulary:",(cnt/tot_cnt)*100)
print("Total Coverage of rare words:",(freq/tot_freq)*100)
     % of rare words in vocabulary: 62.23902087832973
     Total Coverage of rare words: 3.79966494955589
#prepare a tokenizer for reviews on training data
x tokenizer = Tokenizer(num words=tot cnt-cnt)
x_tokenizer.fit_on_texts(list(x_tr))
#convert text sequences into integer sequences
x_tr_seq
                x_tokenizer.texts_to_sequences(x_tr)
x val seq
               x_tokenizer.texts_to_sequences(x_val)
           =
#padding zero upto maximum lengath
            pad sequences(x tr seq,maxlen=max text len, padding='post')
x tr
            pad_sequences(x_val_seq,maxlen=max_text_len, padding='post')
x val
#size of vocabulary ( +1 for padding token)
x_voc = x_tokenizer.num_words + 1
```

```
#prepare a tokenizer for reviews on training data
y tokenizer = Tokenizer()
y_tokenizer.fit_on_texts(list(y_tr))
thresh=7
cnt=0
tot cnt=0
frea=0
tot_freq=0
for key,value in y tokenizer.word counts.items():
    tot cnt=tot cnt+1
    tot freq=tot freq+value
    if(value<thresh):</pre>
        cnt=cnt+1
        freq=freq+value
print("% of rare words in vocabulary:",(cnt/tot cnt)*100)
print("Total Coverage of rare words:",(freq/tot_freq)*100)
     % of rare words in vocabulary: 84.35897435897436
     Total Coverage of rare words: 19.54113081932631
#prepare a tokenizer for reviews on training data
y tokenizer = Tokenizer(num words=tot cnt-cnt)
y_tokenizer.fit_on_texts(list(y_tr))
#convert text sequences into integer sequences
                y tokenizer.texts to sequences(y tr)
y_tr_seq
                y_tokenizer.texts_to_sequences(y_val)
y_val_seq
#padding zero upto maximum length
            pad_sequences(y_tr_seq, maxlen=max_summary_len, padding='post')
y tr
            pad sequences(y val seq, maxlen=max summary len, padding='post')
y val
#size of vocabulary
y_voc = y_tokenizer.num_words +1
y_tokenizer.word_counts['sostok'],len(y_tr)
     (7929, 7929)
ind=[]
for i in range(len(y tr)):
    cnt=0
    for j in y_tr[i]:
        if j!=0:
            cnt=cnt+1
```

```
if(cnt==2):
    ind.append(i)

y_tr=np.delete(y_tr,ind, axis=0)
x_tr=np.delete(x_tr,ind, axis=0)

ind=[]
for i in range(len(y_val)):
    cnt=0
    for j in y_val[i]:
        if j!=0:
            cnt=cnt+1
    if(cnt==2):
        ind.append(i)

y_val=np.delete(y_val,ind, axis=0)
x_val=np.delete(x_val,ind, axis=0)
```

▼ NLP MODEL

```
from tensorflow.keras import layers
import tensorflow as tf
from tensorflow.python.keras import backend as K
logger = tf.get_logger()
class AttentionLayer(tf.keras.layers.Layer):
   This class implements Bahdanau attention (https://arxiv.org/pdf/1409.0473.pdf).
   There are three sets of weights introduced W a, U a, and V a
   def init (self, **kwargs):
        super(AttentionLayer, self).__init__(**kwargs)
   def build(self, input shape):
        assert isinstance(input shape, list)
        # Create a trainable weight variable for this layer.
        self.W a = self.add weight(name='W a',
                                   shape=tf.TensorShape((input shape[0][2], input shape[0][2]
                                   initializer='uniform',
                                   trainable=True)
        self.U_a = self.add_weight(name='U_a',
                                   shape=tf.TensorShape((input shape[1][2], input shape[0][2]
                                   initializer='uniform',
                                   trainable=True)
```

```
self.V a = self.add weight(name='V a',
                               shape=tf.TensorShape((input shape[0][2], 1)),
                               initializer='uniform',
                               trainable=True)
    super(AttentionLayer, self).build(input shape) # Be sure to call this at the end
def call(self, inputs):
    inputs: [encoder_output_sequence, decoder_output_sequence]
    assert type(inputs) == list
    encoder out seq, decoder out seq = inputs
    logger.debug(f"encoder out seq.shape = {encoder out seq.shape}")
    logger.debug(f"decoder out seq.shape = {decoder out seq.shape}")
    def energy step(inputs, states):
        """ Step function for computing energy for a single decoder state
        inputs: (batchsize * 1 * de_in_dim)
        states: (batchsize * 1 * de latent dim)
        logger.debug("Running energy computation step")
        if not isinstance(states, (list, tuple)):
            raise TypeError(f"States must be an iterable. Got {states} of type {type(stat
        encoder full seq = states[-1]
        """ Computing S.Wa where S=[s0, s1, ..., si]"""
        # <= batch size * en seq len * latent dim
        W_a_dot_s = K.dot(encoder_full_seq, self.W_a)
        """ Computing hj.Ua """
        U a dot h = K.expand dims(K.dot(inputs, self.U a), 1) # <= batch size, 1, latent
        logger.debug(f"U a dot h.shape = {U a dot h.shape}")
        """ tanh(S.Wa + hj.Ua) """
        # <= batch size*en seq len, latent dim
        Ws plus Uh = K.tanh(W a dot s + U a dot h)
        logger.debug(f"Ws plus Uh.shape = {Ws plus Uh.shape}")
        """ softmax(va.tanh(S.Wa + hj.Ua)) """
        # <= batch size, en seq len
        e i = K.squeeze(K.dot(Ws plus Uh, self.V a), axis=-1)
        # <= batch size, en seq len
        e i = K.softmax(e i)
```

```
logger.debug(f"ei.shape = {e i.shape}")
            return e_i, [e_i]
        def context_step(inputs, states):
            """ Step function for computing ci using ei """
            logger.debug("Running attention vector computation step")
            if not isinstance(states, (list, tuple)):
                raise TypeError(f"States must be an iterable. Got {states} of type {type(stat
            encoder full seq = states[-1]
            # <= batch_size, hidden_size</pre>
            c i = K.sum(encoder full seq * K.expand dims(inputs, -1), axis=1)
            logger.debug(f"ci.shape = {c i.shape}")
            return c_i, [c_i]
       # we don't maintain states between steps when computing attention
        # attention is stateless, so we're passing a fake state for RNN step function
        fake state c = K.sum(encoder out seq, axis=1)
        fake_state_e = K.sum(encoder_out_seq, axis=2) # <= (batch_size, enc_seq_len, latent_</pre>
        """ Computing energy outputs """
        # e_outputs => (batch_size, de_seq_len, en_seq_len)
        last out, e outputs, = K.rnn(
            energy_step, decoder_out_seq, [fake_state_e], constants=[encoder_out_seq]
        )
        """ Computing context vectors """
        last_out, c_outputs, _ = K.rnn(
            context_step, e_outputs, [fake_state_c], constants=[encoder_out_seq]
        )
        return c_outputs, e_outputs
   def compute_output_shape(self, input_shape):
        """ Outputs produced by the layer """
        return [
            tf.TensorShape((input_shape[1][0], input_shape[1][1], input_shape[1][2])),
            tf.TensorShape((input_shape[1][0], input_shape[1][1], input_shape[0][1]))
        1
from keras import backend as K
K.clear_session()
latent_dim = 150
embedding dim=50
```

```
# Encoder
encoder inputs = Input(shape=(max text len,))
#embedding layer
enc emb = Embedding(x voc, embedding dim,trainable=True)(encoder inputs)
#encoder 1stm 1
encoder lstm1 = LSTM(latent dim,return sequences=True,return state=True,dropout=0.4,recurrent
encoder_output1, state_h1, state_c1 = encoder_lstm1(enc_emb)
#encoder 1stm 2
encoder lstm2 = LSTM(latent dim,return sequences=True,return state=True,dropout=0.4,recurrent
encoder output2, state h2, state c2 = encoder lstm2(encoder output1)
#encoder 1stm 3
encoder_lstm3=LSTM(latent_dim, return_state=True, return_sequences=True,dropout=0.4,recurrent
encoder outputs, state h, state c= encoder lstm3(encoder output2)
# Set up the decoder, using `encoder_states` as initial state.
decoder inputs = Input(shape=(None,))
#embedding layer
dec_emb_layer = Embedding(y_voc, embedding_dim,trainable=True)
dec_emb = dec_emb_layer(decoder_inputs)
decoder_lstm = LSTM(latent_dim, return_sequences=True, return_state=True, dropout=0.4, recurren
decoder outputs, decoder fwd state, decoder back state = decoder lstm(dec emb, initial state=[s
# Attention layer
attn layer = AttentionLayer(name='attention layer')
attn out, attn states = attn layer([encoder outputs, decoder outputs])
# Concat attention input and decoder LSTM output
decoder concat input = Concatenate(axis=-1, name='concat layer')([decoder outputs, attn out])
#dense layer
decoder dense = TimeDistributed(Dense(y voc, activation='softmax'))
decoder_outputs = decoder_dense(decoder_concat_input)
# Define the model
model = Model([encoder inputs, decoder inputs], decoder outputs)
model.summary()
```

Model: "model"

Layer (type)	Output Shape	Param #	Connected to
input_1 (InputLayer)	[(None, 100)]	0	[]
embedding (Embedding)	(None, 100, 50)	734350	['input_1[0][0]']

```
[(None, 100, 150),
 1stm (LSTM)
                                                  120600
                                                             ['embedding[0][0]']
                               (None, 150),
                               (None, 150)]
 input 2 (InputLayer)
                              [(None, None)]
                                                             []
                              [(None, 100, 150),
 1stm 1 (LSTM)
                                                  180600
                                                             ['lstm[0][0]']
                               (None, 150),
                               (None, 150)]
 embedding 1 (Embedding)
                              (None, None, 50)
                                                             ['input 2[0][0]']
                                                  103750
 1stm 2 (LSTM)
                              [(None, 100, 150),
                                                             ['lstm 1[0][0]']
                                                  180600
                               (None, 150),
                               (None, 150)]
 1stm 3 (LSTM)
                              [(None, None, 150),
                                                              ['embedding_1[0][0]',
                                                  120600
                               (None, 150),
                                                               'lstm_2[0][1]',
                               (None, 150)]
                                                               'lstm 2[0][2]']
 attention layer (AttentionLaye
                               ((None, None, 150),
                                                              ['lstm 2[0][0]',
                                                   45150
 r)
                               (None, None, 100))
                                                               'lstm 3[0][0]']
 concat layer (Concatenate)
                              (None, None, 300)
                                                              ['lstm 3[0][0]',
                                                               'attention layer[0][0
 time distributed (TimeDistribu (None, None, 2075) 624575
                                                             ['concat layer[0][0]']
 ted)
______
Total params: 2,110,225
Trainable params: 2,110,225
Non-trainable params: 0
```

#converts the integer sequence to a one-hot vector using sparse_categorical_crossentropy
model.compile(optimizer='rmsprop', loss='sparse categorical crossentropy')

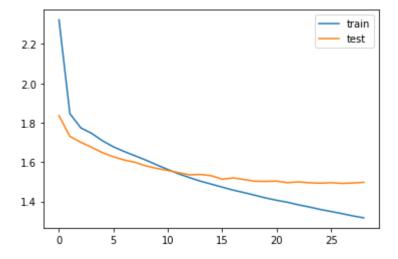
```
es = EarlyStopping(monitor='val_loss', mode='min', verbose=1,patience=2)
```

#Model Training
history=model.fit([x_tr,y_tr[:,:-1]], y_tr.reshape(y_tr.shape[0],y_tr.shape[1], 1)[:,1:] ,epo

```
בע /כ וב
62/62 [============== ] - 360s 6s/step - loss: 1.7083 - val loss: 1.64
Epoch 6/50
62/62 [============== ] - 361s 6s/step - loss: 1.6778 - val loss: 1.62
Epoch 7/50
Epoch 8/50
62/62 [============= ] - 357s 6s/step - loss: 1.6321 - val loss: 1.59
Epoch 9/50
62/62 [============== ] - 362s 6s/step - loss: 1.6101 - val loss: 1.58
Epoch 10/50
62/62 [=============== ] - 359s 6s/step - loss: 1.5858 - val loss: 1.56
Epoch 11/50
Epoch 12/50
62/62 [============== ] - 364s 6s/step - loss: 1.5403 - val loss: 1.54
Epoch 13/50
62/62 [============== ] - 359s 6s/step - loss: 1.5213 - val loss: 1.53
Epoch 14/50
Epoch 15/50
Epoch 16/50
62/62 [============== ] - 360s 6s/step - loss: 1.4731 - val loss: 1.51
Epoch 17/50
Epoch 18/50
62/62 [============== ] - 359s 6s/step - loss: 1.4454 - val loss: 1.51
Epoch 19/50
Epoch 20/50
Epoch 21/50
62/62 [============== ] - 359s 6s/step - loss: 1.4064 - val loss: 1.50
Epoch 22/50
62/62 [============== ] - 365s 6s/step - loss: 1.3964 - val loss: 1.49
Epoch 23/50
62/62 [============ ] - 364s 6s/step - loss: 1.3832 - val_loss: 1.49
Epoch 24/50
Epoch 25/50
Epoch 26/50
Epoch 27/50
Epoch 28/50
62/62 [=============== ] - 359s 6s/step - loss: 1.3271 - val loss: 1.49
Epoch 29/50
```

▼ PLOT OF TRAINING & TESTING

```
from matplotlib import pyplot
pyplot.plot(history.history['loss'], label='train')
pyplot.plot(history.history['val_loss'], label='test')
pyplot.legend()
pyplot.show()
```



reverse_target_word_index=y_tokenizer.index_word
reverse_source_word_index=x_tokenizer.index_word
target word index=y tokenizer.word index

```
# Encode the input sequence to get the feature vector
encoder_model = Model(inputs=encoder_inputs,outputs=[encoder_outputs, state_h, state_c])
# Decoder setup
# Below tensors will hold the states of the previous time step
decoder state input h = Input(shape=(latent dim,))
decoder_state_input_c = Input(shape=(latent_dim,))
decoder hidden state input = Input(shape=(max text len,latent dim))
# Get the embeddings of the decoder sequence
dec emb2= dec emb layer(decoder inputs)
# To predict the next word in the sequence, set the initial states to the states from the pre
decoder outputs2, state h2, state c2 = decoder lstm(dec emb2, initial state=[decoder state in
#attention inference
attn_out_inf, attn_states_inf = attn_layer([decoder_hidden_state_input, decoder_outputs2])
decoder_inf_concat = Concatenate(axis=-1, name='concat')([decoder_outputs2, attn_out_inf])
# A dense softmax layer to generate prob dist. over the target vocabulary
decoder_outputs2 = decoder_dense(decoder_inf_concat)
# Final decoder model
decoder model = Model(
    [decoder_inputs] + [decoder_hidden_state_input,decoder_state_input_h, decoder_state_input_
```

[decoder_outputs2] + [state_h2, state_c2])

```
def decode sequence(input seq):
   # Encode the input as state vectors.
   e out, e h, e c = encoder model.predict(input seq)
   # Generate empty target sequence of length 1.
   target_seq = np.zeros((1,1))
   # Populate the first word of target sequence with the start word.
   target seq[0, 0] = target word index['sostok']
   stop condition = False
   decoded sentence = ''
   while not stop_condition:
        output tokens, h, c = decoder model.predict([target seq] + [e out, e h, e c])
        # Sample a token
        sampled_token_index = np.argmax(output_tokens[0, -1, :])
        sampled token = reverse target word index[sampled token index]
        if(sampled token!='eostok'):
            decoded_sentence += ' '+sampled_token
        # Exit condition: either hit max length or find stop word.
        if (sampled_token == 'eostok' or len(decoded_sentence.split()) >= (max_summary_len-1
            stop condition = True
       # Update the target sequence (of length 1).
        target seq = np.zeros((1,1))
       target_seq[0, 0] = sampled_token_index
        # Update internal states
        e_h, e_c = h, c
   return decoded sentence
def seq2summary(input seq):
   newString=''
   for i in input seq:
        if((i!=0 and i!=target_word_index['sostok']) and i!=target_word_index['eostok']):
            newString=newString+reverse target word index[i]+' '
    return newString
def seq2text(input seq):
   newString=''
   for i in input seq:
        if(i!=0):
            newString=newString+reverse source word index[i]+' '
   return newString
```

→ PREDICTION

```
predicted = []
title =[]
for i in range(0,100):
  print("Abstract:",seq2text(x tr[i]))
  print("Original Title:",seq2summary(y_tr[i]))
  print("Predicted Title:",decode sequence(x tr[i].reshape(1, max text len)))
  print("\n")
  predicted.append(decode_sequence(x_tr[i].reshape(1, max_text_len)))
  title.append(seq2summary(y tr[i]))
   Abstract: models revealed housing insecurity experiences predicted poorer self rated
   Original Title: longitudinal analysis relationship housing insecurity physical health
   1/1 [=======] - 0s 152ms/step
   1/1 [======= ] - 0s 45ms/step
   1/1 [======= ] - 0s 39ms/step
   1/1 [======] - 0s 56ms/step
   1/1 [======= ] - 0s 39ms/step
   1/1 [======= ] - 0s 38ms/step
   1/1 [======= ] - 0s 25ms/step
   Predicted Title: mental health covid pandemic among adults
   1/1 [======= ] - 0s 25ms/step
   1/1 [======= ] - 0s 28ms/step
   1/1 [======= ] - 0s 28ms/step
   1/1 [======= ] - 0s 28ms/step
   1/1 [======= ] - 0s 23ms/step
   1/1 [======] - 0s 24ms/step
   1/1 [======= ] - 0s 26ms/step
   Abstract: dataset finally standardized tests reference intervals obtained compared pu
   Original Title: determination using outpatient data
   1/1 [======] - 0s 27ms/step
   1/1 [=======] - 0s 26ms/step
   Predicted Title: impact covid pandemic
   1/1 [======= ] - 0s 75ms/step
   1/1 [======] - 0s 24ms/step
   1/1 [======] - 0s 25ms/step
   1/1 [======= ] - 0s 28ms/step
   1/1 [=======] - 0s 26ms/step
   Abstract: jet injectors administer assessing impact jet injection pharmaceutical phys
   Original Title: free exploring potential unexpected
   1/1 [=======] - 0s 80ms/step
```

▼ ROUGE SCORE

```
! pip install rouge
     Looking in indexes: <a href="https://pypi.org/simple">https://us-python.pkg.dev/colab-wheels/publications</a>
     Collecting rouge
       Downloading rouge-1.0.1-py3-none-any.whl (13 kB)
     Requirement already satisfied: six in /usr/local/lib/python3.7/dist-packages (from rouge
     Installing collected packages: rouge
     Successfully installed rouge-1.0.1
from rouge import Rouge
rouge = Rouge()
rouge.get scores(title, predicted,avg=True, ignore empty=True)
     {'rouge-1': {'r': 0.230249999999999995,
       'p': 0.10339693639693637,
       'f': 0.1381351230556726},
      'rouge-2': {'r': 0.09092857142857143,
       'p': 0.036539682539682546,
       'f': 0.04997186876638159},
      'rouge-l': {'r': 0.2179166666666662,
       'p': 0.097043401043401,
       'f': 0.12980358531237018}}
res = pd.DataFrame(predicted)
res.columns = ['predicted']
```

res.to_csv("/content/drive/MyDrive/Directed/prediction_result1.csv")

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