### In [1]:

```
# Common imports
import pandas as pd
from IPython.display import Markdown, display, clear_output
```

### In [2]:

```
import _pickle as cPickle
from pathlib import Path
def dumpPickle(fileName, content):
    pickleFile = open(fileName, 'wb')
    cPickle.dump(content, pickleFile, -1)
    pickleFile.close()
def loadPickle(fileName):
    file = open(fileName, 'rb')
    content = cPickle.load(file)
    file.close()
    return content
def pickleExists(fileName):
    file = Path(fileName)
    if file.is file():
        return True
    return False
```

### In [3]:

```
import spacy
from spacy import displacy
nlp = spacy.load('en core web sm')
#Extract answers and the sentence they are in
def extractAnswers(gas, doc):
    answers = []
    senStart = 0
    senId = 0
    for sentence in doc.sents:
        senLen = len(sentence.text)
        for answer in qas:
            answerStart = answer['answers'][0]['answer start']
            if (answerStart >= senStart and answerStart < (senStart + senLen)):</pre>
                answers.append({'sentenceId': senId, 'text': answer['answers'][0
]['text']})
        senStart += senLen
        senId += 1
    return answers
#TODO - Clean answers from stopwords?
def tokenIsAnswer(token, sentenceId, answers):
    for i in range(len(answers)):
        if (answers[i]['sentenceId'] == sentenceId):
            if (answers[i]['text'] == token):
                return True
    return False
#Save named entities start points
def getNEStartIndexs(doc):
    neStarts = {}
    for ne in doc.ents:
        neStarts[ne.start] = ne
    return neStarts
def getSentenceStartIndexes(doc):
    senStarts = []
    for sentence in doc.sents:
        senStarts.append(sentence[0].i)
    return senStarts
def getSentenceForWordPosition(wordPos, senStarts):
    for i in range(1, len(senStarts)):
        if (wordPos < senStarts[i]):</pre>
            return i - 1
def addWordsForParagrapgh(newWords, text):
    doc = nlp(text)
```

```
neStarts = getNEStartIndexs(doc)
    senStarts = getSentenceStartIndexes(doc)
    #index of word in spacy doc text
    i = 0
    while (i < len(doc)):</pre>
        #If the token is a start of a Named Entity, add it and push to index to
 end of the NE
        if (i in neStarts):
            word = neStarts[i]
            #add word
            currentSentence = getSentenceForWordPosition(word.start, senStarts)
            wordLen = word.end - word.start
            shape = ''
            for wordIndex in range(word.start, word.end):
                shape += (' ' + doc[wordIndex].shape )
            newWords.append([word.text,
                             currentSentence.
                            wordLen,
                            word.label ,
                            None,
                            None,
                            None,
                            shapel)
            i = neStarts[i].end - 1
        #If not a NE, add the word if it's not a stopword or a non-alpha (not re
qular letters)
        else:
            if (doc[i].is stop == False and doc[i].is alpha == True):
                word = doc[i]
                currentSentence = getSentenceForWordPosition(i, senStarts)
                wordLen = 1
                newWords.append([word.text,
                                 0,
                                 0,
                                 currentSentence,
                                 wordLen,
                                 None,
                                 word.pos_,
                                 word.tag_,
                                 word.dep ,
                                 word.shape ])
        i += 1
def oneHotEncodeColumns(df):
    columnsToEncode = ['NER', 'POS', "TAG", 'DEP']
    for column in columnsToEncode:
        one hot = pd.get dummies(df[column])
        one_hot = one_hot.add_prefix(column + ' ')
        df = df.drop(column, axis = 1)
        df = df.join(one hot)
    return df
```

```
/home/aayusi/anaconda3/lib/python3.7/site-packages/requests/__init_
_.py:91: RequestsDependencyWarning: urllib3 (1.25.10) or chardet (3.
0.4) doesn't match a supported version!
   RequestsDependencyWarning)
```

### In [4]:

```
def generateDf(text):
    words = []
    addWordsForParagrapgh(words, text)

    wordColums = ['text', 'titleId', 'paragrapghId', 'sentenceId', 'wordCount',
    'NER', 'POS', 'TAG', 'DEP', 'shape']
    df = pd.DataFrame(words, columns=wordColums)

    return df
```

### In [5]:

```
def prepareDf(df):
     #One-hot encoding
     wordsDf = oneHotEncodeColumns(df)
     #Drop unused columns
     columnsToDrop = ['text', 'titleId', 'paragrapghId', 'sentenceId', 'shape']
     wordsDf = wordsDf.drop(columnsToDrop, axis = 1)
     #Add missing colums
     predictorColumns = ['wordCount','NER_CARDINAL','NER_DATE','NER_EVENT','NER_F
AC', 'NER GPE', 'NER LANGUAGE', 'NER LAW', 'NER LOC', 'NER MONEY', 'NER NORP', 'NER ORD
INAL', 'NER ORG', 'NER PERCENT', 'NER_PERSON', 'NER_PRODUCT', 'NER_QUANTITY', 'NER_TIM
E','NER_WORK_OF_ART','POS_ADJ','POS_ADP','POS_ADV','POS_CCONJ','POS_DET','POS_IN
TJ','POS_NOUN','POS_NUM','POS_PART','POS_PRON','POS_PROPN','POS_PUNCT','POS_SYM','POS_VERB','POS_X','TAG_''','TAG_-LRB-','TAG_.','TAG_ADD','TAG_AFX','TAG_CC','TAG_CD','TAG_DT','TAG_EX','TAG_FW','TAG_IN','TAG_JJ','TAG_JJR','TAG_JJS','TAG_LS'
,'TAG_MD','TAG_NFP','TAG_NN','TAG_NNP','TAG_NNPS','TAG_NNS','TAG_PDT','TAG_POS',
'TAG_PRP','TAG_PRP$','TAG_RB','TAG_RBR','TAG_RBS','TAG_RP','TAG_SYM','TAG_TO','T
AG_UH', 'TAG_VB', 'TAG_VBD', 'TAG_VBG', 'TAG_VBN', 'TAG_VBP', 'TAG_VBZ', 'TAG_WDT', 'TAG
_WP','TAG_WRB','TAG_XX','DEP_ROOT','DEP_acl','DEP_acomp','DEP_advcl','DEP_advmod','DEP_amod','DEP_amod','DEP_attr','DEP_aux','DEP_auxpass','DEP_ca
se','DEP_cc','DEP_ccomp','DEP_compound','DEP_conj','DEP_csubj','DEP_csubjpass',
'DEP_dative','DEP_dep','DEP_det','DEP_dobj','DEP_expl','DEP_intj','DEP_mark','DEP_meta','DEP_neg','DEP_nmod','DEP_npadvmod','DEP_nsubj','DEP_nsubjpass','DEP_num mod','DEP_oprd','DEP_parataxis','DEP_pcomp','DEP_pobj','DEP_poss','DEP_preconj',
'DEP predet', 'DEP prep', 'DEP prt', 'DEP punct', 'DEP quantmod', 'DEP relcl', 'DEP xc
omp'l
     for feature in predictorColumns:
           if feature not in wordsDf.columns:
                 wordsDf[feature] = 0
      return wordsDf
```

### In [16]:

```
def predictWords(wordsDf, df):
    predictorPickleName = 'nb-predictor.pkl'
    predictor = loadPickle(predictorPickleName)

    y_pred = predictor.predict_proba(wordsDf)

    labeledAnswers = []
    for i in range(len(y_pred)):
        labeledAnswers.append({'word': df.iloc[i]['text'], 'prob': y_pred[i][0]]})

    return labeledAnswers
```

### In [17]:

```
def blankAnswer(firstTokenIndex, lastTokenIndex, sentStart, sentEnd, doc):
    leftPartStart = doc[sentStart].idx
    leftPartEnd = doc[firstTokenIndex].idx
    rightPartStart = doc[lastTokenIndex].idx + len(doc[lastTokenIndex])
    rightPartEnd = doc[sentEnd - 1].idx + len(doc[sentEnd - 1])

    question = doc.text[leftPartStart:leftPartEnd] + '____' + doc.text[rightPartStart:rightPartEnd]
    return question
```

### In [18]:

```
def addQuestions(answers, text):
    doc = nlp(text)
    currAnswerIndex = 0
    qaPair = []
    #Check wheter each token is the next answer
    for sent in doc.sents:
        for token in sent:
            #If all the answers have been found, stop looking
            if currAnswerIndex >= len(answers):
                break
            #In the case where the answer is consisted of more than one token, c
heck the following tokens as well.
            answerDoc = nlp(answers[currAnswerIndex]['word'])
            answerIsFound = True
            for j in range(len(answerDoc)):
                if token.i + j >= len(doc) or doc[token.i + j].text != answerDoc
[j].text:
                    answerIsFound = False
            #If the current token is corresponding with the answer, add it
            if answerIsFound:
                question = blankAnswer(token.i, token.i + len(answerDoc) - 1, se
nt.start, sent.end, doc)
                gaPair.append({'question' : question, 'answer': answers[currAnsw
erIndex]['word'], 'prob': answers[currAnswerIndex]['prob']})
                currAnswerIndex += 1
    return qaPair
```

### In [19]:

```
def sortAnswers(qaPairs):
    orderedQaPairs = sorted(qaPairs, key=lambda qaPair: qaPair['prob'])
    return orderedQaPairs
```

## In [20]:

```
import gensim
from gensim.test.utils import datapath, get_tmpfile
from gensim.models import KeyedVectors

glove_file = 'glove.6B.300d.txt'
tmp_file = 'word2vec-glove.6B.300d.txt'

from gensim.scripts.glove2word2vec import glove2word2vec
glove2word2vec(glove_file, tmp_file)
model = KeyedVectors.load_word2vec_format(tmp_file)
```

### In [24]:

```
def generate_distractors(answer, count):
    answer = str.lower(answer)

##Extracting closest words for the answer.
    try:
        closestWords = model.most_similar(positive=[answer], topn=count)
    except:
        #In case the word is not in the vocabulary, or other problem not loading
    embeddings
        return []

#Return count many distractors
    distractors = list(map(lambda x: x[0], closestWords))[0:count]
    return distractors
```

### In [25]:

```
def addDistractors(qaPairs, count):
    for qaPair in qaPairs:
        distractors = generate_distractors(qaPair['answer'], count)
        qaPair['distractors'] = distractors

return qaPairs
```

### In [26]:

```
def generateQuestions(text, count):
    # Extract words
    df = generateDf(text)
    wordsDf = prepareDf(df)
    # Predict
    labeledAnswers = predictWords(wordsDf, df)
    # Transform questions
    gaPairs = addQuestions(labeledAnswers, text)
    # Pick the best questions
    orderedQaPairs = sortAnswers(gaPairs)
    # Generate distractors
    questions = addDistractors(orderedQaPairs[:count], 4)
    # Print
    for i in range(count):
        display(Markdown('### Question ' + str(i + 1) + ':'))
        print(questions[i]['question'])
        display(Markdown('#### Answer:'))
        print(questions[i]['answer'])
        display(Markdown('#### Incorrect answers:'))
        for distractor in questions[i]['distractors']:
            print(distractor)
        print()
```

### In [55]:

text1 = "Bansoori is an Indian classical instrument. Akhil plays Bansoori and Gu itar. Puliyogare is a South Indian dish made of rice and tamarind. Priya writes poems. Osmosis is the movement of a solvent across a semipermeable membrane tow ard a higher concentration of solute. In biological systems, the solvent is typi cally water, but osmosis can occur in other liquids, supercritical liquids, and even gases. I like Bansoori. When a cell is submerged in water, the water molecules pass through the cell membrane from an area of low solute concentration to high solute concentration. For example, if the cell is submerged in saltwater, water molecules move out of the cell. If a cell is submerged in freshwater, wat er molecules move into the cell. Raja-Yoga is divided into eight steps, the firs t is Yama non killing, truthfulness, non stealing, continence, and non receiving of any gifts."

# In [62]:

generateQuestions(text1, 10)

/home/aayusi/anaconda3/lib/python3.7/site-packages/sklearn/base.py:3
34: UserWarning: Trying to unpickle estimator GaussianNB from versio
n 0.20.3 when using version 0.23.2. This might lead to breaking code
or invalid results. Use at your own risk.
UserWarning)

Qı	uesti	ion 1	•							
Ιt	was	July	21,	1969,	and	Neil	Armstrong	 with	а	start.

#### Answer:

awoke

### **Incorrect answers:**

woke awakened awakens awaken

# **Question 2:**

The journey had begun several days earlier, when on July 16th, the A pollo 11 \_\_\_\_\_ from Earth headed into outer space.

### Answer:

launched

### **Incorrect answers:**

launch
launching
launches
initiated

# **Question 3:**

The journey had begun several days earlier, when on July 16th, the A pollo 11 launched from Earth \_\_\_\_ into outer space.

### Answer:

headed

### **Incorrect answers:**

heading heads head arrived

# **Question 4:**

Take out all of the trash, and all of the dirty dishes in the kitchen sink.
Answer:
place
Incorrect answers:
places time
where
Question 5:
For any remaining items, see if you can squeeze them in under your bed or them into the back of your closet.
Answer:
stuff
Incorrect answers:
things
thing
really guys
Question 6:
Oceans and have much in common, but they are also quite differ ent.
Answer:
lakes
Incorrect answers: lake
rivers
ponds streams
Question 7:
Both have plants and living in them.
Answer:
animals
Incorrect answers:

animal humans birds mammals

# **Question 8:**

When it is time for a vacation, both will make a great place to visi t and \_\_\_\_\_.

### Answer:

enjoy

### **Incorrect answers:**

enjoying enjoyed enjoys want

# **Question 9:**

The journey had begun several days earlier, when on July 16th, the \_ 11 launched from Earth headed into outer space.

### Answer:

Apollo

### **Incorrect answers:**

spacecraft
gemini
astronauts
moon

# **Question 10:**

\_\_\_\_ are usually surrounded by land, while oceans are what surround continents.

### Answer:

Lakes

### **Incorrect answers:**

lake rivers ponds streams

```
In [60]:
```

```
f = open("file1.txt", 'r')
text1 = f.read()
```

# In [61]:

generateQuestions(text1, 10)

/home/aayusi/anaconda3/lib/python3.7/site-packages/sklearn/base.py:3 34: UserWarning: Trying to unpickle estimator GaussianNB from versio n 0.20.3 when using version 0.23.2. This might lead to breaking code or invalid results. Use at your own risk.

UserWarning)

Question	1:
----------	----

It was July 21, 1969, and Neil Armstrong \_\_\_\_ with a start.

#### Answer:

awoke

#### Incorrect answers:

woke awakened awakens awaken

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The journey had begun several days earlier, when on July 16th, the A pollo 11 from Earth headed into outer space.

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heading heads head arrived

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Take out all of the trash, and all of the dirty dishes in the kitchen sink.
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place
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where
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For any remaining items, see if you can squeeze them in under your bed or them into the back of your closet.
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In a sum of an arrange
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animals
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0	u	е	S	ti	O	n	8:
τ		_	_		_		_

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## Incorrect answers:

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1	in [ ]:
1	in [ ]: