CSF 256 FA24: NLP UCSD PA3:

Retrieval-Augmented Generation (RAG) (40 points)

The goal of this assignment is to gain hands-on experience with aspects of **Retrieval-Augmented Generation (RAG)**, with a focus on retrieval. You will use **LangChain**, a framework that simplifies integrating external knowledge into generation tasks by:

- Implementing various vector databases for efficient neural retrieval. You will use a vector database for storing our memories.
- Allowing seamless integration of pretrained text encoders, which you will access via HuggingFace models. You will use a text encoder to get text embeddings for storing in the vector database.

Data

You will build a retrieval system using the <u>QMSum Dataset</u>, a human-annotated benchmark designed for question answering on long meeting transcripts. The dataset includes over 230 meetings across multiple domains.

Release Date: November 6, 2024 | Due Date: November 18, 2024

IMPORTANT: After copying this notebook to your Google Drive along with the two data files, paste a link to your copy below. To create a publicly accessible link:

- 1. Click the Share button in the top-right corner.
- 2. Select "Get shareable link" and copy the link.

Link: paste your link here:

https://colab.research.google.com/drive/1vF6PICLMZLKxkkiFmXERME-_7boUh1nE?usp=sharing

Notes:

Make sure to save the notebook as you go along.

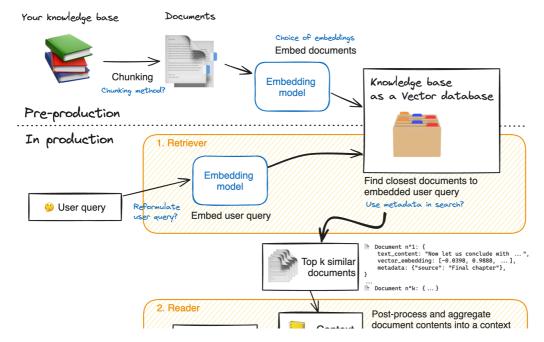
Submission instructions are located at the bottom of the notebook.

```
# This mounts your Google Drive to the Colab VM.
from google.colab import drive
drive.mount('/content/drive')
# TODO: Enter the foldername in your Drive where you have saved this notebook
# e.g. 'CSE156/assignments/PA3/'
FOLDERNAME = 'CSE256_PA3'
assert FOLDERNAME is not None, "[!] Enter the foldername."
# Now that we've mounted your Drive, this ensures that
# the Python interpreter of the Colab VM can load
# python files from within it.
import sys
sys.\,path.\,append\,('\,\underline{/content/drive/My}\quad Drive/\,\{\}\,'.\,format\,(FOLDERNAME)\,)
# This is later used to use the IMDB reviews
%cd /content/drive/My\ Drive/$FOLDERNAME/
    Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive", force_remount=True).
     /content/drive/My Drive/CSE256_PA3
```

RAG Workflow

Retrieval-Augmented Generation (RAG) systems involve several interconnected components. Below is a RAG workflow diagram from Hugging Face. Areas highlighted in blue indicate opportunities for system improvement.

In this assignment, we will focus on the *Retriever so the PA does not cover any processes starting from "2. Reader" and below.



First, install the required model dependancies.

```
pip install -q torch transformers langchain_chroma bitsandbytes langchain faiss-gpu langchain_huggingface langchain-community sentence-tra
    from \quad tqdm. \, notebook \quad import \quad tqdm
import pandas as pd
import os
import csv
import sys
import numby
           as np
import time
import random
from typing import Optional, List, Tuple
import matplotlib.pyplot as plt
import textwrap
import torch
seed = 42
random. seed (seed)
np. random. seed (seed)
torch.manual_seed(seed)
torch.cuda.manual seed all(seed)
torch.backends.cudnn.deterministic = True
torch.backends.cudnn.benchmark = False
# Disable huffingface tokenizers parallelism
os.environ["TOKENIZERS_PARALLELISM"] = "false"
```

Load the meetings dataset

```
from langchain.docstore.document import Document

def load_documents(doc_file):
    """
    Loads the document contents from the first file.

    :param doc_file: Path to the document file (document ID <TAB> document contents).
    :return: A dictionary {document_id: document_contents}.
    """
    max_size = sys.maxsize
    csv.field_size_limit(max_size)

    documents = {}
    with open(doc_file, 'r', encoding='utf-8') as f:
        reader = csv.reader(f, delimiter='\t')
        for row in reader:
```

Retriever - Building the retriever

The retriever functions like a search engine: given a user query, it returns relevant documents from the knowledge base.

These documents are then used by the Reader model to generate an answer. In this assignment, however, we are only focusing on the retriever, not the Reader model.

Our goal: Given a user question, find the most relevant documents from the knowledge base.

Key parameters:

- top_k: The number of documents to retrieve. Increasing top_k can improve the chances of retrieving relevant content.
- chunk size: The length of each document. While this can vary, avoid overly long documents, as too many tokens can overwhelm most reader models.

Langchain offers a huge variety of options for vector databases and allows us to keep document metadata throughout the processing.

1. Specify an Embedding Model and Visualize Document Lengths

🛬 /usr/local/lib/python3.10/dist-packages/sentence_transformers/cross_encoder/CrossEncoder.py:13: TqdmExperimentalWarning: Using `tqdm.autonotebool from tadm, autonotebook import tadm, trange

/usr/local/lib/python3.10/dist-packages/huggingface_hub/utils/_token.py:89: UserWarning:

The secret `HF_TOKEN` does not exist in your Colab secrets.

To authenticate with the Hugging Face Hub, create a token in your settings tab (https://huggingface.co/settings/tokens), set it as secret in your You will be able to reuse this secret in all of your notebooks.

Please note that authentication is recommended but still optional to access public models or datasets.

warnings.warn(

tokenizer config.json: 100%

385/385 [00:00<00:00, 25.9kB/s] modules.json: 100% README.md: 100% 68.1k/68.1k [00:00<00:00, 3.70MB/s]

sentence_bert_config.json: 100% 57.0/57.0 [00:00<00:00, 4.70kB/s]

583/583 [00:00<00:00, 26.3kB/s] config.json: 100%

model.safetensors: 100% 66.7M/66.7M [00:00<00:00, 185MB/s] 394/394 [00:00<00:00, 10.5kB/s]

vocab.txt: 100% 232k/232k [00:00<00:00, 6.72MB/s]

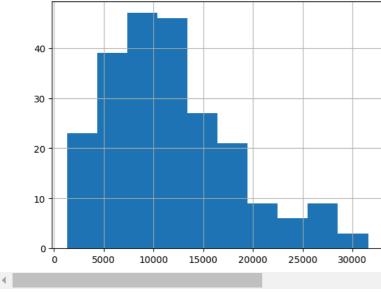
712k/712k [00:00<00:00, 4.39MB/s] tokenizer.ison: 100%

special tokens map.json: 100% 125/125 [00:00<00:00, 2.97kB/s] 1_Pooling/config.json: 100% 190/190 [00:00<00:00, 4.88kB/s]

Model's maximum sequence length: 512

100% 230/230 [00:13<00:00, 31.36it/s]

Distribution of document lengths in the knowledge base (in count of tokens)



2. Split the Documents into Chunks

The documents (meeting transcripts) are very long—some up to 30,000 tokens! To make retrieval effective, we'll split each document into smaller, semantically meaningful chunks. These chunks will serve as the snippets the retriever compares to the query, returning the top_k most relevant ones.

Objective: Create Semantically Relevant Snippets

Chunks should be long enough to capture complete ideas but not so lengthy that they lose focus.

We will use Langchain's implementation of recursive chunking with ${\tt RecursiveCharacterTextSplitter}.$

- Parameter chunk_size controls the length of individual chunks: this length is counted by default as the number of characters in the chunk.
- Parameter chunk_overlap lets adjacent chunks get a bit of overlap on each other. This reduces the probability that an idea could be cut in half by the split between two adjacent chunks.

From the produced plot below, you can see that now the chunk length distribution looks better!

```
from \quad langehain.\ text\_splitter \quad import \quad Recursive Character Text Splitter
text_splitter = RecursiveCharacterTextSplitter(
        chunk size = 768,
        chunk_overlap = 128.
doc_snippets = text_splitter.split_documents(docs)
print(f''Total \ \{len(doc\_snippets)\} \ snippets \ to \ be \ stored \ in \ our \ vector \ store.")
```

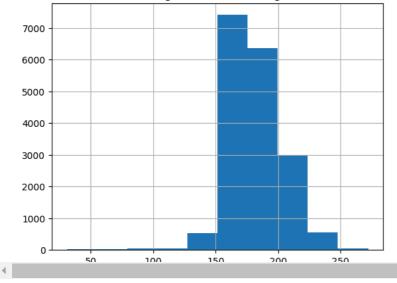
```
lengths = [len(tokenizer.encode(doc.page_content)) for doc in tqdm(doc_snippets)]

# Plot the distribution of document snippet lengths, counted as the number of tokens
fig = pd.Series(lengths).hist()
plt.title("Distribution of document lengths in the knowledge base (in count of tokens)")
plt.show()
```

Total 18070 snippets to be stored in our vector store.

00% 18070/18070 [00:11<00:00, 1074.26it/s]





3. Build the Vector Database

To enable retrieval, we need to compute embeddings for all chunks in our knowledge base. These embeddings will then be stored in a vector database.

How Retrieval Works

A query is embedded using an embedding model and a similarity search finds the closest matching chunks in the vector database.

The following cell builds the vector database consisting of all chunks in our knowledge base.

```
from \ langehain\_hugging face \ import \ Hugging Face Embeddings
from langchain.vectorstores import FAISS
from langchain_community.vectorstores.utils import DistanceStrategy
# Automatically set the device to 'cuda' if available, otherwise use 'cpu'
device = "cuda" if torch.cuda.is_available() else "cpu"
print(f"Found device: {device}")
embedding_model = HuggingFaceEmbeddings(
       model name=EMBEDDING MODEL NAME,
        multi_process=True,
        model_kwargs={"device": device},
        encode kwargs={"normalize embeddings": True}, # Set True for cosine similarity
start_time = time.time()
\label{eq:knowledge_vector_database} \mbox{ = } \mbox{ FAISS.from\_documents} \mbox{ (}
        \verb|doc_snippets|, & \verb|embedding_model|, & \verb|distance_strategy=DistanceStrategy|. COSINE| \\
end_time = time.time()
elapsed\_time = (end\_time - start\_time)/60
print(f"Time taken: {elapsed_time} minutes")
    Found device: cuda
     Time taken: 1.4466180642445883 minutes
```

4. Querying the Vector Database

Using LangChain's vector database, the function <code>vector_database.similarity_search(query)</code> implements a Bi-Encoder (covered in class), independently encoding the query and each document into a single-vector representation, allowing document embeddings to be precomputed.

Let's define the Bi-Encoder ranking function and then use it on a sample query from the QMSum dataset.

```
## The function for ranking documents given a query:
def rank_documents_biencoder(user_query, top_k = 5):
                       Function for document ranking based on the query.
                       :param query: The query to retrieve documents for.
                       :return: A list of document IDs ranked based on the query (mocked).
                       retrieved_docs = KNOWLEDGE_VECTOR_DATABASE.similarity_search(query=user_query, k=top_k)
                       for i. doc in enumerate(retrieved docs):
                                               ranked list.append(retrieved docs[i].metadata['source'])
                       return ranked_list # ranked document IDs.
{\tt user\_query} \ = \ {\tt "what} \ {\tt did} \ {\tt kirsty} \ {\tt williams} \ {\tt am} \ {\tt say} \ {\tt about} \ {\tt her} \ {\tt plan} \ {\tt for} \ {\tt quality} \ {\tt assurance} \ ?{\tt "matchings} \ {\tt matchings} \ {\tt matchings} \ {\tt plan} \ {\tt for} \ {\tt quality} \ {\tt assurance} \ {\tt matchings} \ {\tt m
retrieved docs = rank documents biencoder(user query)
print("\n======Top-5 documents======="")
print("\n\nRetrieved documents:", retrieved_docs)
print("\n=
 \overline{\Sigma}
                  Retrieved documents: ['doc_211', 'doc_2', 'doc_43', 'doc_160', 'doc_43']
```

5. TODO: Implementation of ColBERT as a Reranker for a Bi-Encoder (35 points)

The Bi-Encoder's ranking for the sample query is not optimal: the ground truth document is not ranked at position 1, instead the document ID, doc_211 is ranked at position 1. To determine the correct document ID for this query, refer to the questions_answers. tsv file.

In this task, you will implement the <u>ColBERT</u> approach by Khattab and Zaharia. We'll use a simplified version of ColBERT, focusing on the following key steps:

- 1. Retrieve the top (K = 15) documents for query (q) using the Bi-Encoder.
- 2. Re-rank these top (K = 15) documents using ColBERT's fine-grained interaction scoring. This will involve:
 - Using frozen BERT embeddings from a HuggingFace BERT model (no training is required, thus our version is not expected to work as well as full-fledged ColBERT).
 - o Calculating scores based on fine-grained token-level interactions between the query and each document.
- 3. Implement the method rank_documents_finegrained_interactions() to perform this re-ranking.
 - o Test your method on the same query as in the cell from #4 above.
 - Print out the entire re-ranked document list of 5 document IDs, as done in #4 above (the code below does it for you)
- 4. Ensure that your ColBERT implementation ranks the correct document at position 1 for the sample query.

Note: Since the same document is divided into multiple chunks that retain the original document ID, you may see the same document ID appear multiple times in your top_k results. However, each instance refers to a different chunk of the document's content.

```
Returns:
      - ranked list of document IDs.
      \mbox{\tt\#} Step 1: Retrieve the top-K documents
      retrieved_docs = KNOWLEDGE_VECTOR_DATABASE.similarity_search(query=user_query, k=shortlist)
      # Tokenize and embed the query
      query_inputs = tokenizer(user_query, return_tensors='pt', truncation=True, padding=True)
      with torch.no_grad():
             # Step 2: Calculate fine-grained interaction scores for each document
      doc scores = []
      for doc in retrieved docs:
             # Tokenize and embed the document
             doc_inputs = tokenizer(doc.page_content, return_tensors='pt', truncation=True, padding=True)
             with torch.no_grad():
                   # Step 3: Compute fine-grained token interaction scores
             # Cosine similarity between each query token and each document token
             similarity_scores = F.cosine_similarity(query_embeddings.unsqueeze(2), doc_embeddings.unsqueeze(1), dim=-1)
             # Aggregate scores by taking the max similarity for each query token
                                  = similarity scores.max(dim=2)
             max similarity scores,
             document_score = max_similarity_scores.sum().item()
             # Store document ID and its score
             doc_scores.append((doc.metadata['source'], document_score))
      \mbox{\tt\#} Step 4: Sort documents by score and retrieve top-k
      ranked\_docs = sorted(doc\_scores, key=lambda \ x: \ x[1], \ reverse=True) \ [:top\_k]
      ranked\_list = [doc\_id \ for \ doc\_id, \ \_ \ in \ ranked\_docs]
      return ranked list # ranked document IDs
user query = "what did kirsty williams am say about her plan for quality assurance ?"
retrieved_docs = rank_documents_finegrained_interactions(user_query)
print("\n=====Top-5 documents======
print("\n\nRetrieved documents:", retrieved_docs)
print("\n=
\rightarrow
            -----Top-5 documents-----
    Retrieved documents: ['doc 2', 'doc 160', 'doc 43', 'doc 2', 'doc 43']
```

6. TODO: ColBERT Max vs. Mean Pooling for Relevance Scoring of Documents: (5 points)

ColBERT uses a form of **max pooling**, where each query term's contribution to the relevance score of a document is determined by its maximum similarity to any document term. One alternative approach is **mean pooling**, where each query term's contribution is calculated as the average similarity across all document terms.

Discuss the merits and potential limitations of using mean pooling versus max pooling in ColBERT. In your answer, consider how each approach might affect retrieval accuracy, sensitivity to specific token matches, interpretability, and anything you deem relevant. You are welcome to use an analysis of ColBERT's performance on the provided sample query in your discussion, but this is not required.

Explain here (<= 5 sentences):

Max pooling in ColBERT emphasizes key term matches by focusing on the highest similarity for each query term, improving precision and sensitivity to important terms but risking over-sensitivity to isolated matches. In contrast, mean pooling offers a balanced relevance score by averaging similarities, which captures broader contextual matches and improves recall but may dilute the impact of highly relevant terms. Max pooling is generally preferred in ColBERT to prioritize exact matches, especially for precise or entity-specific queries, while mean pooling could be more suitable for general or descriptive queries.

Submission Instructions

- 1. Click the Save button at the top of the Jupyter Notebook.
- 2. Select Runtime -> Run All. This will run all the cells in order, and will take several minutes.
- 3. Once you've rerun everything, save a PDF version of your notebook. Make sure all your code and answers are displayed in the pdf, it's okay if the provided codes get cut off because lines are not wrapped in code cells).
- 4. Look at the PDF file and make sure all your code and answers are there, displayed correctly. The PDF is the only thing your graders will see!
- 5. Submit your PDF on Gradescope.

7. (Optional) Full evaluation pipeline for your own exploration.

For this assignment, we only ask you to explore one sample query. Running on many queries is super slow without the right compute. If you have compute/and/or time to wait, below is a more complete evaluation setup that works with all the queries in QMSum dataset, and reports the precision@k=5 metric.

Note: you need to remove the comment markers from the code below.

```
#
  def load_questions_answers(qa_file):
#
          Loads the questions and corresponding ground truth document IDs.
#
          :param ga file: Path to the question-answer file (document ID <TAB> question <TAB> answer).
#
          :return: A list of tuples [(document_id, question, answer)].
#
          qa_pairs = []
          with open(qa_file, 'r', encoding='utf-8') as f:
#
                 reader = csv.reader(f, delimiter='\t')
Ħ
                  for row in reader:
#
                         doc_id, question, answer = row
#
                         qa_pairs.append((doc_id, question, answer))
#
          random.shuffle(qa_pairs)
#
          return qa pairs
#
  def precision_at_k(ground_truth, retrieved_docs, k):
#
          Computes Precision at k for a single query.
#
          :param ground_truth: The name of the ground truth document.
Ħ
          :param retrieved_docs: The list of document names returned by the model in ranked order.
          :param k: The cutoff for computing Precision.
#
          :return: Precision at k.
Ħ
#
          return 1 if ground truth in retrieved docs[:k] else 0
#
  def evaluate(doc_file, qa_file, ranking_fuction = None, k= 5):
#
#
          Evaluate the retrieval system based on the documents and question-answer pairs.
#
          :param doc_file: Path to the document file.
#
                 ga file: Path to the question-answer file.
          :param k: The cutoff for Precision@k.
#
#
          # Load the QA pairs
#
          qa_pairs = load_questions_answers(qa_file)
          precision_scores = []
#
          for doc_id, question, _ in qa_pairs:
                  retrieved_docs = ranking_fuction(question)
#
                  precision_scores.append(precision_at_k(doc_id, retrieved_docs, k))
                  avg_precision_at_k = sum(precision_scores) / len(precision_scores)
#
                  if len(precision_scores) %10==0:
                         print(f'' After \quad \{len(precision\_scores)\} \quad queries, \quad Precision@\{k\}: \quad \{avg\_precision\_at\_k\}'')
#
          # Compute average Precision@k
          avg\_precision\_at\_k \ = \ sum(precision\_scores) \ / \ len(precision\_scores)
```

```
# print(f"Precision@{k}: {avg_precision_at_k}")

# qa_file = 'questions_answers.tsv'  # document ID <TAB> question <TAB> answer

# start_time = time.time()
# evaluate(doc_file, qa_file,rank_documents_biencoder)
# end_time = time.time()
# elapsed_time = (end_time - start_time)/60
```