```
In [ ]:
In [1]: import os, pickle, requests, pdfplumber
        from dotenv import load_dotenv
        from langchain.schema import Document
        from langchain.embeddings import OpenAIEmbeddings
        from langchain.vectorstores import FAISS
        from langchain.llms import OpenAI
        from langchain.chains import RetrievalQA
        import gradio as gr
        load_dotenv()
        API KEY = os.getenv("OPENAI API KEY")
        assert API_KEY, "Please set OPENAI_API_KEY in your .env file"
       C:\Users\joshu\anaconda3\envs\rag-env\lib\site-packages\tqdm\auto.py:21: TqdmWarnin
       g: IProgress not found. Please update jupyter and ipywidgets. See https://ipywidget
       s.readthedocs.io/en/stable/user_install.html
         from .autonotebook import tqdm as notebook_tqdm
In [ ]:
In [2]: # Cell 2 (simplified): Download one EMA PDF & convert to text
        import os, requests, pdfplumber
        os.makedirs("data", exist_ok=True)
        # Only EMA Bioanalytical Method Validation
        url = "https://www.ema.europa.eu/en/documents/scientific-guideline/guideline-bioana
        r = requests.get(url, timeout=30)
        r.raise_for_status()
        pdf_path = "data/ema_bioanalytical_guideline.pdf"
        with open(pdf_path, "wb") as f:
            f.write(r.content)
        # Convert EMA PDF -> .txt
        txt_path = pdf_path.replace(".pdf", ".txt")
        pages = []
        with pdfplumber.open(pdf_path) as pdf:
            for page in pdf.pages:
                pages.append(page.extract_text() or "")
        with open(txt_path, "w", encoding="utf-8") as f:
            f.write("\n\n".join(pages))
        print(" ☑ EMA guideline downloaded and converted to text in ./data/")
       EMA guideline downloaded and converted to text in ./data/
In [ ]:
```

```
In [ ]:
In [3]: from langchain.schema import Document
                   import os
                   def load_docs(folder="data"):
                            docs = []
                            for fn in os.listdir(folder):
                                      if not fn.endswith(".txt"):
                                               continue
                                     with open(os.path.join(folder, fn), encoding="utf-8") as f:
                                               txt = f.read()
                                      docs.append(Document(page_content=txt, metadata={"source": fn}))
                            return docs
                   all_docs = load_docs("data")
                   print(f" \( \) Loaded \( \
                 Loaded 1 text docs for indexing.
In [ ]:
In [ ]:
In [4]: # Cell 4: Build & persist FAISS index with a local HF embedder
                   !pip install --quiet sentence-transformers
                   from langchain.embeddings import HuggingFaceEmbeddings
                   from langchain.vectorstores import FAISS
                   import pickle
                   embedder = HuggingFaceEmbeddings(model_name="all-MiniLM-L6-v2")
                   faiss_index = FAISS.from_documents(all_docs, embedder)
                   with open("faiss_index.pkl", "wb") as f:
                            pickle.dump(faiss_index, f)
                   print("☑ FAISS index built locally (no OpenAI calls) and saved to faiss_index.pkl'
                C:\Users\joshu\AppData\Local\Temp\ipykernel_3512\3554573147.py:8: LangChainDeprecati
                onWarning: The class `HuggingFaceEmbeddings` was deprecated in LangChain 0.2.2 and w
                ill be removed in 1.0. An updated version of the class exists in the :class:`~langch
                ain-huggingface package and should be used instead. To use it run `pip install -U :c
                lass:`~langchain-huggingface` and import as `from :class:`~langchain_huggingface imp
                ort HuggingFaceEmbeddings``.
                     embedder = HuggingFaceEmbeddings(model_name="all-MiniLM-L6-v2")
                 FAISS index built locally (no OpenAI calls) and saved to faiss_index.pkl
```

```
In [ ]:
In [ ]:
In [5]: # Cell 5: Wire up the RAG chain with GPT-3.5 using the v1.0+ API
        from langchain.chat_models import ChatOpenAI
        from langchain.chains import RetrievalQA
        # (If you restarted, reload the FAISS store:)
        # with open("faiss_index.pkl", "rb") as f:
              faiss_index = pickle.load(f)
        11m = ChatOpenAI(
            model_name="gpt-3.5-turbo",
            temperature=0,
            openai_api_key=API_KEY
        retriever = faiss_index.as_retriever(search_kwargs={"k": 5})
        rag = RetrievalQA.from_chain_type(
            11m=11m,
            chain_type="stuff",
            retriever=retriever,
            return_source_documents=True
        print(" RAG chain is ready!")
       C:\Users\joshu\AppData\Local\Temp\ipykernel_3512\774492991.py:10: LangChainDeprecati
       onWarning: The class `ChatOpenAI` was deprecated in LangChain 0.0.10 and will be rem
       oved in 1.0. An updated version of the class exists in the :class:`~langchain-openai
       package and should be used instead. To use it run `pip install -U :class:`~langchain
       -openai` and import as `from :class:`~langchain_openai import ChatOpenAI``.
         11m = ChatOpenAI(
       RAG chain is ready!
In [ ]:
In [6]: def answer_with_sources(question):
            res = rag({"query": question})
            answer = res["result"]
            sources = "\n".join(f"- {d.metadata['source']}" for d in res["source_documents"
```

```
return f"**Answer:**\n{answer}\n\n**Sources:**\n{sources}"

iface = gr.Interface(
    fn=answer_with_sources,
    inputs=gr.Textbox(lines=2, placeholder="Ask medical questions..."),
    outputs="markdown",
    title="% Medical RAG Demo",
    description="GPT-3.5 + EMA Bioanalytical Guideline"
)

iface.launch(share=True)
```

- * Running on local URL: http://127.0.0.1:7860
- * Running on public URL: https://c73119fb403c441137.gradio.live

This share link expires in 1 week. For free permanent hosting and GPU upgrades, run `gradio deploy` from the terminal in the working directory to deploy to Hugging Face Spaces (https://huggingface.co/spaces)

Medical RAG Demo

GPT-3.5 + EMA Bioanalytical Guideline

question

What parameters and acceptance criteria does the EMA Bioanalytical Method Validation guideline recommend for accuracy and precision?

Clear Submit

Answer: The EMA Bioanalytical Method Validation guideline recommends the following parameters and acceptance criteria for accuracy and precision:

1. Accuracy:

- Within-run accuracy: The mean concentration should be within 15% of the nominal values for the QC samples, except for the LLOQ which should be within 20%.
- Between-run accuracy: The mean concentration should be within 15% of the nominal values for the QC samples, except for the LLOQ which should be within 20%.

2. Precision:

• Within-run precision: The coefficient of variation (CV) value should not exceed 15% for the QC samples, except for the LLOQ which should not exceed 20%.

Out[6]:

```
C:\Users\joshu\AppData\Local\Temp\ipykernel_3512\3418708132.py:2: LangChainDeprecati
onWarning: The method `Chain.__call__` was deprecated in langchain 0.1.0 and will be
removed in 1.0. Use :meth:`~invoke` instead.
  res = rag({"query": question})
```

In []:	
In []:	