





How to hack for a cause

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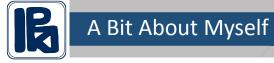








Giorgos Kosta
Research Engineer
CaSToRC
eurocc.cyi.ac.cy



- Research Engineer at CaSToRC
- Awareness and Communications task leader for EuroCC2
- Contact & Info:
 - Email: g.kosta@cyi.ac.cy





What our datasets look like.

What we expect from you

The proposed methodology and tools.



How a patient file looks likes

		-1			
Age	78				
Medicine	Dosage	Frequency	Route	Allergies	Co-existing conditions
Phenergan-Promethazine	25mg	prn , nocte	orally	N/A	Hypokalemia
Colchicine	0.5mg	b.d.	orally		Pulmonary toxicity from chemotherapy
DNS (dextrose & sodium chloride)	500ml	o.d.	iv		acute kidney injury
Human albumin 20g/100ml	2 vials	b.d.	iv		thrombocytopenia
Meropenem	1gr	t.d.s.	iv		Arterial hypertension
Bioflor	1 tablet	t.d.s.	orally		Diabetes
Loperamide	2mg	prn max dose 16mg/24h	orally		dyslipidemia
Magnesium eff.	300mg	b.d.	orally		1 2434.1432.139
Dexamethasone	4mg	o.d.	iv		
Betaloc (metoprolol)	50mg	o.d.	orally		
allopurinol	100mg	o.d.	orally		
Crestor (rosuvastatin)	5mg	nocte	orally		
Paracetamol	1gr	prn / max dosage 4gr/24h	iv		
Metoclopramide	10mg	pro	iv		
Ondasetron	4mg	b.d.	iv or orally		
Pantoprazole	40mg	o.d.	iv or orally		
Kcl - potassium chloride	30 cc	o.d.	iv		
Atrovent 500mg (ipratropium)	1 ampoule	t.d.s.	neb.		
Tramadex / Tramadol	50mg	o.d.	orally		
Pregabalin 50mg	50mg	b.d.	orally		
interactions					
Tramadol + Pregabalin					
KCI + Metoprolol					
Tramadol + Ondansetron					



Starting point for going through medicine files...

```
v medicine files
from langchain community.document loaders import PyPDFLoader
from tgdm import tgdm
                                                                                                 Acyclovir.pdf
directory='medicine files'
                                                                                                 Allopurinol 100mg.pdf
files = [f for f in os.listdir(directory) if f.endswith('.pdf')]
                                                                                                 Amitriptyline 10 mg.pdf
                                                                                                 Apixaban.pdf
all splits = []
                                                                                                 aspirin.pdf
                                                                                                 Atorvastatin.pdf
for file name in tqdm(files):
                                                                                                 Atrovent 500mg (ipratropium).pdf
    file path = os.path.join(directory, file name)
                                                                                                 Betaloc (metoprolol) 50mg.pdf
    # print(f"Processing {file path}...") # Optional: for tracking progress
                                                                                                 Bioflor.pdf
    # Load the PDF
                                                                                                 biperidine.pdf
    loader = PyPDFLoader(file path)
                                                                                                 Bisoprolol.pdf
                                                                                                 Buscopan.pdf
     . . .
                                                                                                 Carbamazepine 200mg.pdf
```

You are free to download any more medicine information from the web!

- Patient files not entirely consistent same medicine names are written in a different way (e.g haloperidole vs haloperidol).
 - You will be given a script to read the patient files though!

 The medicine leaflets that will be provided contain a lot of useless information.



What we expect from you

- A list of all explained interactions per patient found in the patient files. Deadline Sunday 13/10/2024 at 23:59
- The code that generates your the above list. Deadline Sunday 13/10/2024 at 23:59
- A presentation on Monday showcasing and explaining your solution.
- Winner will be announced in the future when we evaluate your results.



Main deliverable

• We expect the major interactions of the whole patient file. For patient 2, it should look something like this:

```
{ "PATIENT 2":
                     "interaction":["tramadol","pregabalin"],
                     "severity":"Major",
                     "Explanation":
                                           "Using narcotic pain medications together with other medications that cause central
                                           nervous system depression can lead to respiratory distress, coma, and even death."
              },
                     "interaction":["X","Y"],
                     "severity":"Minor",
                     "Explanation":
                                           "There were no major interactions found in the dataset,
                                           however for medicine X it is suggested to consult with your doctor."
```

You must use most of the patient information because:

- If you iterate through all combinations of possible interactions, you will run out of time (nCr):
 - Medicine to medicine
 - Medicine to allergy
 - Medicine to co-existing conditions
 - Etc.
- You may generate a lot of false positives because in the medicine leaflets they mention the phrase "Consult with your doctor if ..." a lot!



Solving the problem

1. Read through patient files and extract important information

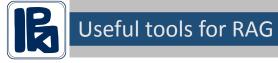
2. Match this information with the appropriate medicine leaflets

Use an LLM to give an answer based on the information you gathered.



How to use RAG to win this Hackathon

USING LOCAL LLMS WITH LOCAL DATA BASE MODEL What was the name of that Sorry, I don't have access to movie Chris emailed me about any specific information about last year? emails or ... BASE MODEL + USER DATASET What was the name of that Chris Thomson emailed you movie Chris emailed me about about "The Fall" on October last year? 23rd last year. The Fall is an adventure fantasy film released in 2006, starring... Library



Langchain - Create chain of prompts

- Langgraph Create sophisticated LLM agents
- Langsmith Track your chains for debugging

Ollama - Your local LLM inference server

1 GPU per team - 32GB V100 from NVIDIA

- Limited available models:
 - Gemma2:27b (Google)
 - Llama3.1:8b (Meta)
 - Phi3.5:3.8b-mini-instruct-fp16 (Microsoft)
 - Nemotron-mini:4b-instruct-fp16 (Nvidia)

Limited context size depending on model



Conclusion and Questions

- Discord server with private channels for each team.
 - If you need any help mention me @costacis21
- Github repository with our slides and more.



Thank you for the attention!

More information:



https://castorc.cyi.ac.cy/



Contact us at:

<u>eurocc-contact@cyi.ac.cy</u>













Funded by the European Union. This work has received funding from the European High Performance Computing Joint Undertaking (JU) and Germany, Bulgaria, Austria, Croatia, Cyprus (co-funded by the EU within the framework of the Cohesion Policy Programme "THALIA 2021-2027"), Czech Republic, Denmark, Estonia, Finland, Greece, Hungary, Ireland, Italy, Lithuania, Latvia, Poland, Portugal, Romania, Slovenia, Spain, Sweden, France, Netherlands, Belgium, Luxembourg, Slovakia, Norway, Türkiye, Republic of North Macedonia, Iceland, Montenegro, Serbia under grant agreement No 101101903.