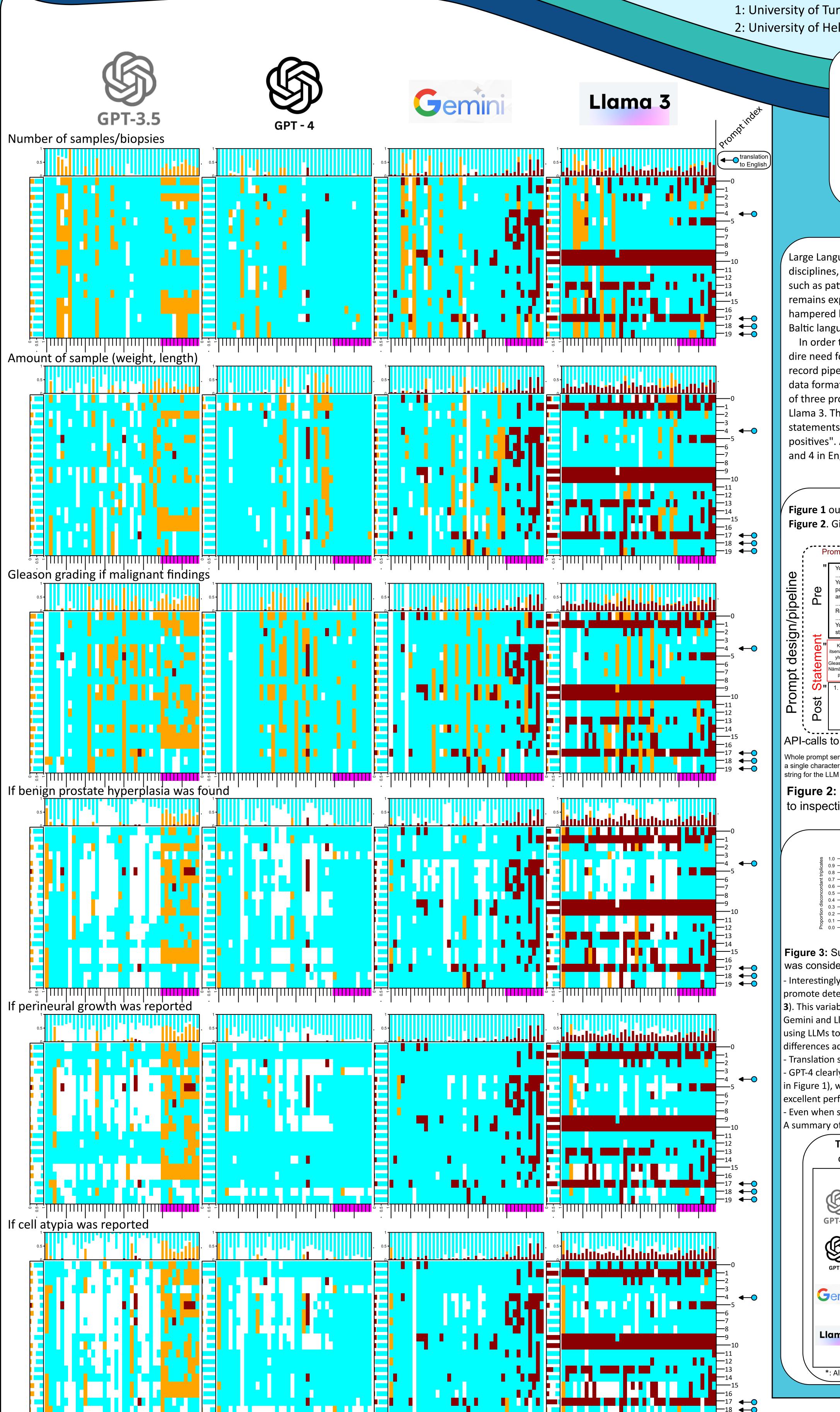
P6

## Practical comparison of GPT, Gemini and Llama LLMs for structuring biomedical data from synthetic Finnish prostate cancer pathology statements

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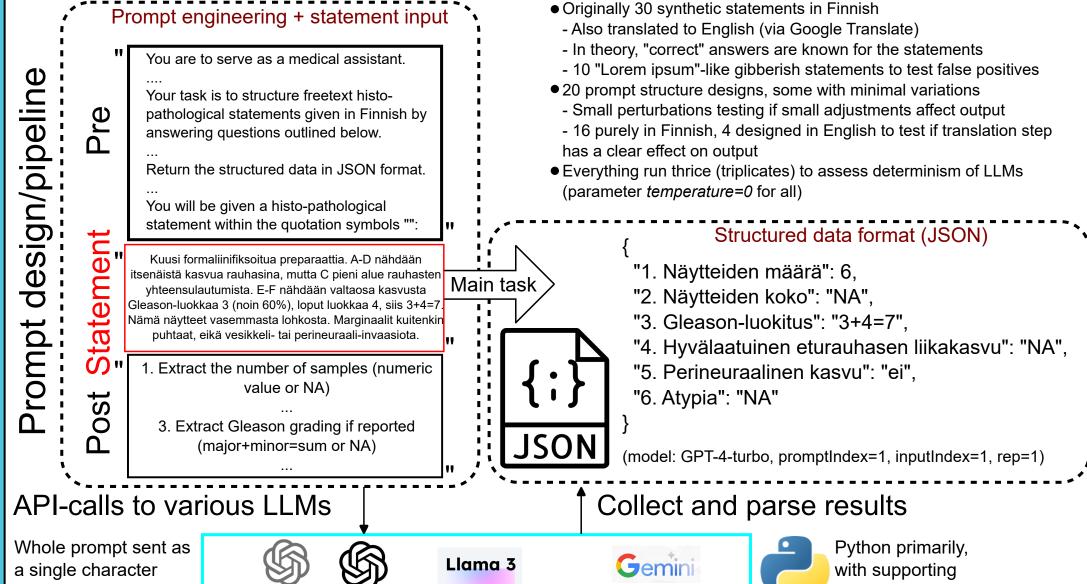
## Introduction

Large Language Models (LLMs) have made significant breakthroughs across all scientific disciplines, including processing of biomedical text data. However, for specific domains such as patient confidentiality retaining electronic health records, usage of such models remains experimental and challenging. Deploying such models to real use cases is further hampered by limited availability of teaching data for smaller languages such as Nordic or Baltic languages.

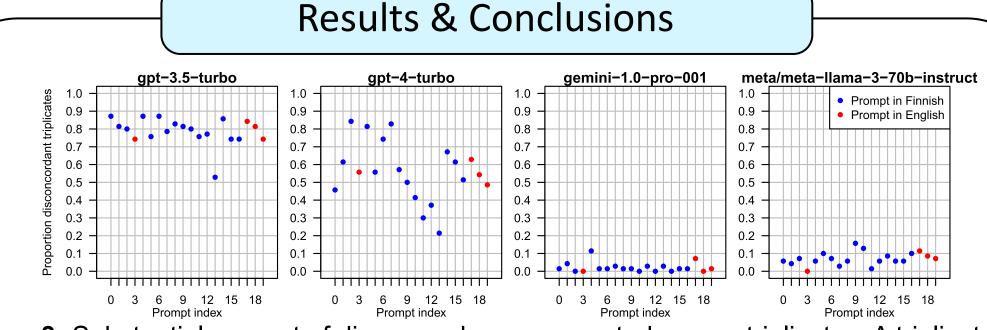
In order to model data for research purposes and e.g. efficient visualizations, there is dire need for structuring freetext fields produced via multitude of electronic health care record pipelines. LLMs offer great potential in structuring freetext via simple structured data formats such as JSON, XML, or even tab-separated values. Here, we examine the use of three prominent LLMs for JSON structuring: GPT (both 3.5 and 4 releases), Gemini, and Llama 3. The utilized data comprises of 30 synthetic Finnish histo-pathological statements, as well as 10 non-sensical input statements testing LLM behaviour for "false positives". A total of 20 prompt designs were used, of which 16 were purely in Finnish and 4 in English coupled with the statements first machine-translated to English.

## Materials & Methods

Figure 1 outlines the whole set of results collected via running a pipeline outlined in Figure 2. GitHub link for running the study: https://github.com/Syksy/HistPath\_LLMpy



**Figure 2:** Rough outline for generating the results from prompt engineering to inspecting the newly structured data.



**Figure 3:** Substantial amount of disconcordance was noted across triplicates. A triplicate was considered disconcordant if even one of the three contained a single diffent character.

- Interestingly, as all runs were conducted as triplicates (with *temperature*-parameter set at 0 to promote determinism), a notable amount of variability was observed across the triplicates (**Figure 3**). This variability was notably highest in the GPT-family (both 3.5 and 4), but larger than zero also in Gemini and Llama. This highlights the fact that if exact scientific reproducibility is to be expected, using LLMs to structure data within a research pipeline poses a risk of producing stochastic differences across runs.

- Translation step to English (prompt indices 4, 17, 18, 19) had very minimal if any impact
- GPT-4 clearly outperformed GPT-3.5, particularly in avoiding "false positives" (pink minor columns in Figure 1), while both GPT-4 and Gemini presented stable performance. Llama 3 presented both excellent performance, but also severe inconsistencies for JSON structure (prompts 1, 9, 10, and 17)
- Even when set to minimal blockage, Gemini's harm/profanity filters prevented model predictions.

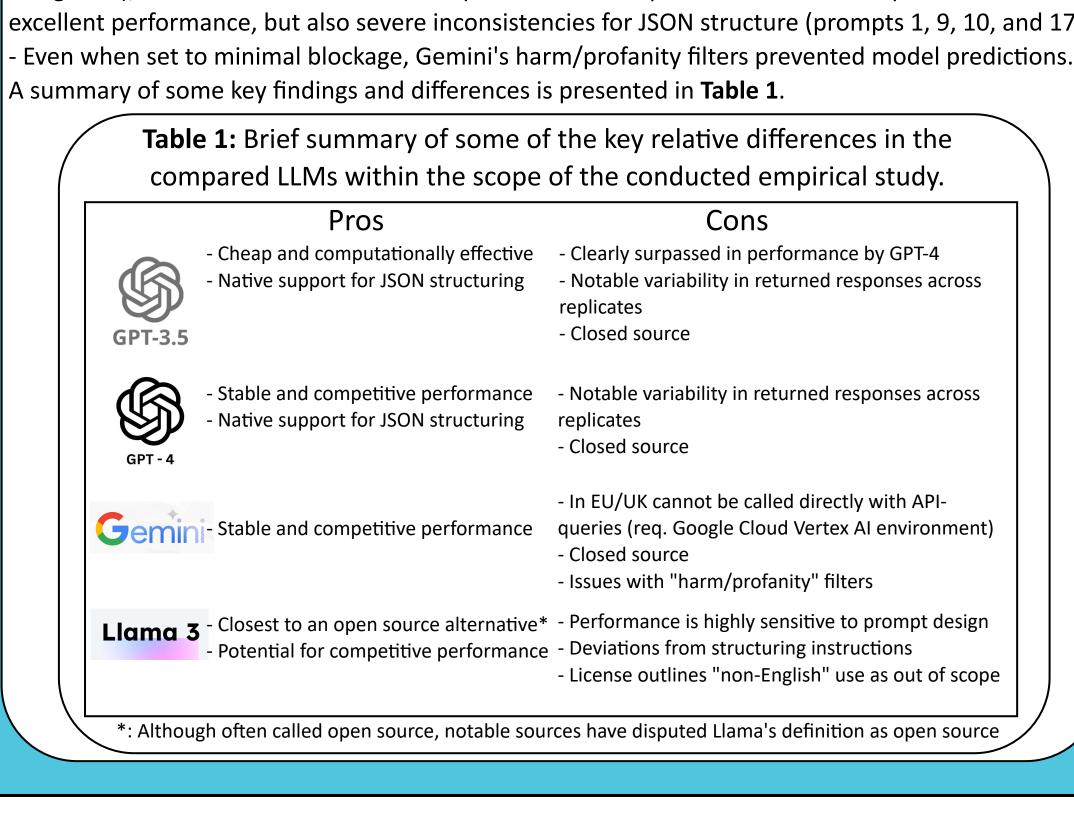


Figure 1: 30 synthetic Finnnish histo-pathological statements along with 10 "false positive" nonsensical statements were given to four distinct LLMs (major columns). 6 questions were systematically queried across all prompt designs (major rows). Minor rows depict 20 different prompt designs, and minor columns the different input statements. Colour codings indicate comparison against a known "correct" answer, or if the query failed. Marginal distributions of answer correctness are shown on top (per statement) and left (per prompt design) in the heatmap cell.

JSON parsing error
Input statements with nonsense or offtopic content

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