

Graph parsing application for bio problems

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Nowadays input data for parsing algorithms are not limited to be linear strings, and context-free grammars are used not only for programming languages specification. One of classical examples is a context-free path querying for graph data bases where input is a graph and path constraints are specified by a grammar.

We have some experience in the areas mentioned above [?, ?]. GLL-based context-free path querying algorithm [?] implemented by the authors is faster than solution which was presented at ISWC-2016 [?]. We have some ideas of graph parsing applications in bio data analysys. For example, context-free pattern search in metagenomical assemblies. We also want to investigate practical areas of application and to create solutions based on our framework to demonstrate its practical value.

References

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