Programming Expertise University of Potsdam SS 2022 Detlef Groth, Christian Kappel Test-Exam July 20^{nh}, 2022

You have 90 minutes time for the implementation of the programming tasks. Make after the steps 1-3 intermediate versions: GooboParser1.cpp, GooboParser2.cpp and GooboParser3.cpp . Submit at the end of the exam those single versions by USB stick or by E-Mail preferentially as zip archive (surname.zip) to me (E-Mail: dgroth@uni-potsdam.de). 75% of the tasks will be on execute correctness and 25% of the tasks will be weighted by usefulness and clearness of the implementation.

The theory questions in 4-6 are to be answered first without any aids and the sheet with the answers is to be handed in after about 15 minutes. For the computer tasks 1-3 all aids are allowed during the exam. This does not include personal support from fellow students or other persons.

Clarification

With this I state, that I will not take and give any not allowed support during the exam.

Name, Matrikel-Number.:

Signature:

- 1. 2 points (layout console application)
- 2. 4 points (implementation of console application)
- 3. 4 points (advanced terminal task)
- 4. 2 points (theory C)
- 5. 2 points (theory C)
- 6. 2 points (theory C++)

Sum: 16 points

Good luck !!

1. Layout of console application and utilizing command line arguments (2 points):

Create the basic outline of a console application with main function, help function and checking of command line arguments. Save the possible command line arguments in variables or use a command line processor like argparse. If the right number of arguments was not given call the help function and exit the application. The two arguments and the optional third should be a goobo input filename and a taskname like *getMetaCyc* or *tabMetaCyc* and a *MetaCyc* identifier for *getMetaCyc*.

You find the obofiles on Moodle. Check if the given filename points to a valid filename, check if a valid command is entered and check, if getMetaCyc was provided, that as well an id argument was given and that this id contains either a RXN or a PWY substring.

C++ filename	first task:	

2. Opening and searching in the Goobo file (4 points): Our program should work with any goobo file. Please don't hardcode the filename in your application, if you do, you get a minus point. Implement the getMetaCyc function for a specific MetaCyc id. If the user gives the for example the three arguments: *filename.obo getMetaCyc BETA-LACTAMASE-RXN* on the command line, then go-obo file is parsed and the id, the name, the namespace and and the entry itself should displayed on the terminal. Your functiomn has to return the data to the function caller, for instance as a vector of tuples, print the result outside of the function, for instance in main.

Hint: If your code is very slow use limit your search to the first 1000 entries first to save your programming time during development. You should uncomment this in your final program. Here a possible example invocation:

GO:0008800	beta-lactamase activity m	iolecular_function	MetaCyc:BETA-L	ACTAMASE-RXN
GO:0033250	penicillinase activity mole	ecular_function	MetaCyc:BETA-LAC	TAMASE-RXN
GO:0033251	cephalosporinase activity	molecular_fun	ction MetaCyc:B	ETA-LACTAMASE-RXN

C++ filename(s) second task:

3. Advanced Terminal Application (4 points):

1 (1 point): Extend your application so that as a C++ class and so that it can handle more than 1 MetaCyc id at the same time given on the terminal. Hint: handle the multiple ids in main, not within the function/method!

2) (3 points): Create a mapping using the command argument MCSummary between all RXN-Metacyc entries, GO-ids and EC ids, only if all three entries are within the same go-id display them, here an example:

\$ appname go-2020.obo tabMetaCyc

GO:0000016 EC:3.2.1.108 LACTASE-RXN

GO:0000034 EC:3.5.4.2 ADENINE-DEAMINASE-RXN GO:0000048 EC:2.3.2.12 PEPTIDYLTRANSFERASE-RXN

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Hint: You can assume, that EC comes first before the MetaCyc id if there is any. In case there is no EC-mapping write NA in the EC column.