  (https://profile.intra.42.fr/searches)

mes-sadk

(https://
profile.intra.42.fr)

SCALE FOR PROJECT OCAML - PATTERN MATCHING AND DATA TYPES - 0 (/PROJECTS/ OCAML-PATTERN-MATCHING-AND-DATA- TYPES-0)

You should evaluate 1 student in this team



Git repository

git@vogosphere-v2.1337.ma:vogosphere/intra-uuid-2fef5b8f-9b7b-49ba



Introduction


For the good of this evaluation, we ask you to:

- Stay mannerly, polite, respectful and constructive during this evaluation. The trust between you and the 42 community depends on it.
- Bring out to the graded student (or team) any mistake she or he might did.
- Accept that there might be differences of interpretation of the subject or the rules between you and the graded student (or team). Stay open minded and grade as honestly as possible.

Guidelines

- You must grade only what is present and the graded student's (or team) repository.
- You must stop grading at the first failed exercise, but you are encouraged to continue testing and discussing the following exercises.

Attachments

 subject.pdf (<https://cdn.intra.42.fr/pdf/pdf/144531/en.subject.pdf>)

Preliminaries

Respect of the rules

- The graded student (or team) work is present on her or his repository.
- The graded student (or team) is able to explain her or his work at any time of the evaluation.
- The general rules and the possible day-specific rules are respected at any time of the evaluation.
- For this project, you need to clone the Git repository on the evaluated person's computer.

☒ Yes

☐ No

OCAML - Pattern Matching and Data Types - 0

- For each exercise, you must compile the exercise using `ocamlc` and run the generated executable. If the compilation fails or warns, or an exception is thrown at runtime, the exercise is failed. - Whether the graded student provided tests or not, you must test her or his work extensively and assess if the work is done or not. - Remember to check function names, TYPES, behaviours and outputs. - Never test overflows for today. - One more time, you HAVE to check the type of the functions since it's essential today

Ex00, Do you even compress bro?

Test the program with at least the following lists: encode
`["a","a","a","b","b","b"]` encode `[]` encode `[(3, 2); (3, 2); (4, 3)]`

☒ Yes

☐ No

Ex01, Crossover

Test the program with at least the following lists : crossover
`[1;2;3] [1;2;3]` crossover `["toto","tata","titi"]`
`["toto","tata","tutu"]` crossover `["toto","tata","titi"] []`
 crossover `[] ["toto","tata","titi"]`

☒ Yes

☐ No

Ex02, Fifty Strings of Gray

Test the Gray sequences generation program with at least the following values : 0, 1, 2, 3, 4 and 5. Also if the student has

used the list concatenation operator (@), It's considered as cheating so -42.

☒ Yes☐ No

Ex03, One and one and one is three

Test the function with the following values. sequence 1 = "1"
sequence -1 = "" sequence 6 = "312211" sequence 8 = "1113213211"

☒ Yes☐ No

Ex04, DNA -> Nucleotides

Check the following facts :

- phosphate and deoxyribose types are both string type aliases.
- nucleobase type is a variant composed of A, T, C, G, and None.
- nucleotide type is a record type or a triplet composed of one of every types above (e.g. 1 phosphate, 1 deoxyribose and 1 nucleobase).

Also check the generate_nucleotide function. It should return a nucleotide type from a char as a parameter.

☒ Yes☐ No

Ex05, DNA -> Helix

- Check the function generate_helix (1pts)
- Check the function helix_to_string (1pts)
- Check the function complementary_helix (for example ATCG should output TAGC) (3pts)

Rate it from 0 (failed) through 5 (excellent)



Ex06, DNA -> Messenger RNA

Check the function generate_rna. The ATCG helix must produce an UAGC rna.

☒ Yes☐ No

Ex07, DNA -> Ribosome

Check the function `generate_base_triplets`. it should return a nucleotide triplet list from an rna as parameter. If the number of nucleotides in the rna is not a factor of 3, the triplet should be forced filled with `None` elements (Or anything that can represent the Great Interstellar Void).

Check the `decode_arn` function. It should return a protein (a list of aminoacids) from an element of type rna as parameter. Use a couple different arn sequences of your choice to test this using the `string_of_protein` function.

☒ Yes

☐ No

Ex08, DNA -> The Complete Process of Protein Creation

This one is a gift, check that the function is functioning as intended. Life is amazing isn't it ? For example the parameter "TACTACATCTAC" should output something like Met-Met-STOP in the end. Also check that each step is explained thoroughly and clearly outputed. Remember that the decoding function **MUST** stop the process at the first STOP occurrence.

☒ Yes

☐ No

Ratings

Don't forget to check the flag corresponding to the defense

☒ Ok

☐ Empty work

☐ Incomplete work

☐ Invalid compilation

☐ Cheat

☐ Crash

☐ Incomplete group

☐ Concerning situation

☐ Forbidden function

☐ Can't support / explain code

Conclusion

Leave a comment on this evaluation

Finish evaluation

Declaration on
the use of
cookies ([https://](https://profile.intra.42.fr/)
profile.intra.42.fr/

General term of
use of the site
([https://](https://profile.intra.42.fr/)
profile.intra.42.fr/

Legal notices
([https://](https://profile.intra.42.fr/legal/terms/3)
[profile.intra.42.fr/](https://profile.intra.42.fr/legal/terms/3)

Privacy policy
([https://](https://profile.intra.42.fr/legal/terms/5)
[profile.intra.42.fr/](https://profile.intra.42.fr/legal/terms/5)

Rules of
procedure
([https://](https://profile.intra.42.fr/)
profile.intra.42.fr/

Terms of use for
video
surveillance
(<https://>

[legal/terms/2\)](#)

[legal/terms/6\)](#)

[legal/terms/4\)](#)

[profile.intra.42.fr/
legal/terms/1\)](#)