MULTIPLE SEQUENCE ALIGNMENT FOR ANTIMICROBIAL RESISTANCE

Team QTON

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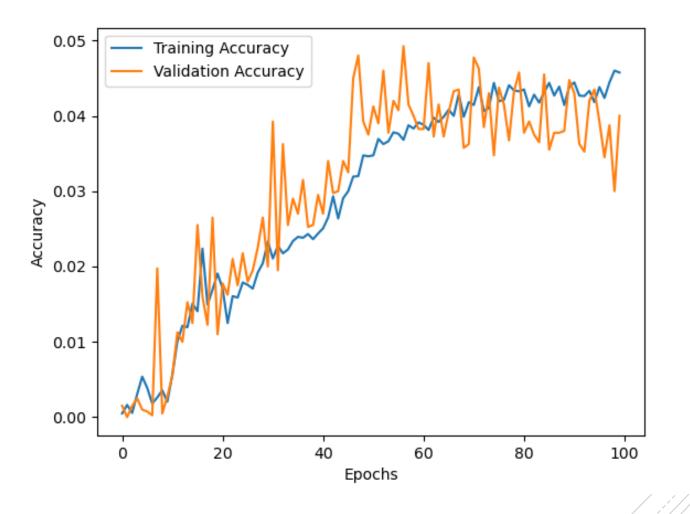




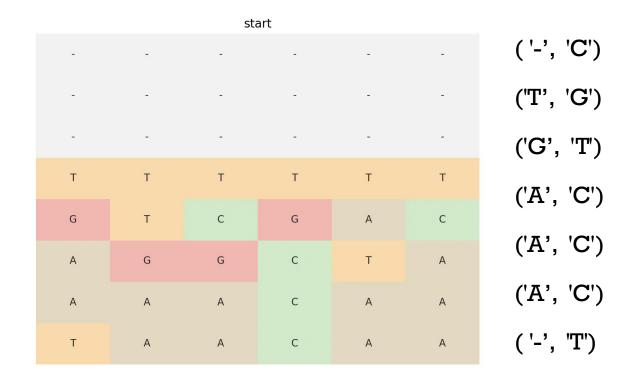
Introduction: MSA for AMR

OBJECTIVE: Train a model to make optimal moves

FCNN Strategy 1



Method Strategy 2 generate the index of valued index for each sequence and each position



Method Strategy 2

repeat for each position, try each no valued sequence

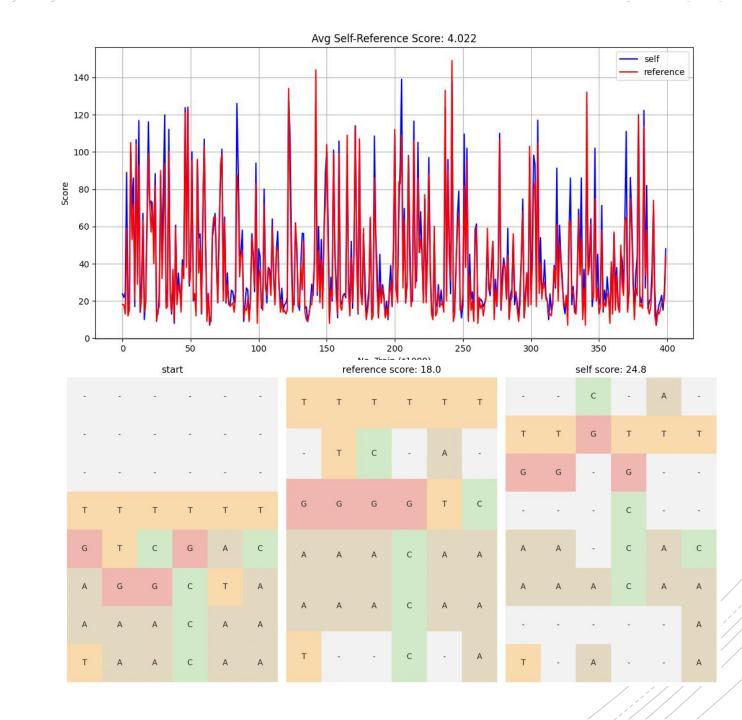
	0	1	2	3	4	5
0	1.00000	0.00000	0.00000	0.00000	0.00000	0.00000
1	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
2	1.00000	0.00000	0.00000	1.00000	0.00000	1.00000
3	0.00000	0.00000	1.00000	0.00000	1.00000	1.00000
4	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
5	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
6	0.00000	0.00000	0.00000	0.00000	0.00000	0.00000
7	1.00000	1.00000	1.00000	1.00000	1.00000	1.00000

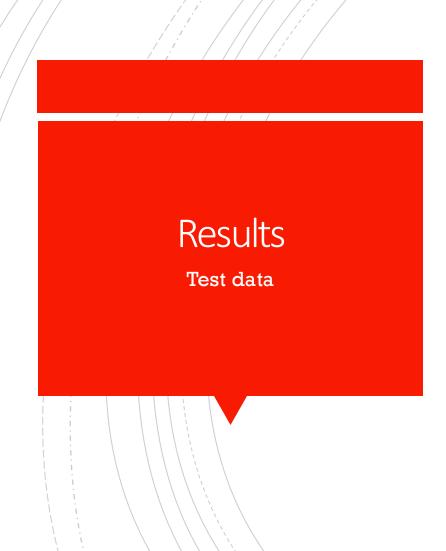
• for example:

- column (position) 0;
- try sequence 1,3,4,5,6,
- calculate the score 9,9,9,9,9
- select maximum sequence 1, select step (2,0), if score increase keep this step, other wise, discard.

Results

Train data: on average, our score higher than reference score





Test.26 Score: 37

Test.27 Score: 61.65

■ Test.28 Score: 123.25

• Test.29 Score: 89

• Test.30 Score: 14

■ Test.31 Score: 12

• Test.32 Score: 42.15

• Test.33 Score: 84

• Test.34 Score: 18.9

■ Test.35 Score: 22.9

Test. 193 Score: 59

Test.194 Score: 59

Test.195 Score: 19.9

Test.196 Score: 118

Test.197 Score: 18.9

Test.198 Score: 19.9

■ Test.199 Score: 69

■ Test.200 Score: 90

■ Test.201 Score: 90

Test.202 Score: 95

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- Data Handling and Feature Engineering
- Model Development
- Puzzle Solving
- Expectations

