

Exercise sheet 6: BLAST

Exercise 1

You are given accession number NM_000667.3. Use the BLAST web server to find out about the gene that belongs to this accession number (choose **nucleotide blast**, and the database **reference RNA sequences (refseq_rna)**).

1a)

Which gene is it, and in which organism?

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Solution Gene: Alcohol Dehydrogenase 1A

Organism: *Homo sapiens* (human)

1b)

Which other organisms does it seem to be highly conserved in?

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Solution

- *Gorilla gorilla*: gorilla
- *Pan troglodytes*: common chimpanzee
- *Pan paniscus*: bonobo
- *Nomascus leucogenys*: northern white-cheeked gibbon
- *Cebus capucinus*: white-headed capuchin

Many more...

Exercise 2

You are given a nucleotide query sequence $q = \text{ATAC}$, and a nucleotide database sequence $s = \text{ATAAAACGGGGG}$. The word-size $k = 2$. Use a simple scoring scheme that assigns a score of 2 for a match and a score of -1 for a mismatch.

2a)

Generate all k -length words of the query sequence.

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Solution

- $w_1 = AT$
- $w_2 = TA$
- $w_3 = AC$

2b)

List all possible words for the first k -length word (AT) that have a score of at least $T_1 = 1$.

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Solution

- $s(AA) = 1$
- $s(AC) = 1$
- $s(AG) = 1$
- $s(AT) = 4$
- $s(CT) = 1$
- $s(GT) = 1$
- $s(TT) = 1$

2c)

Scan the database for exact matches for the words from the question 3B.

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Solution AA at position 2,3,4. AC at position 5, AT at position 0.

2d)

Extend the exact matches that you found in the question 3C to the left/right and report all MSPs with a score greater than 4.

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Solution AA:

```
Pos: 2      ATA
           |||
           AAA      with score 3
```

```
Pos: 3      ATAC
           ||||
           AAAC      with score 5
```

```
Pos: 4      AT
           ||
           AA      with score 1
```

AT:

```
Pos: 0      ATA
           |||
           ATA      with score 6
```

AC:

```
Pos: 5      AT
           ||
           AC      with score 1
```

MSPs start in the template at index 0 and 3.

2e)

What happens if we vary the parameters k and T_1 ?

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Solution

- Higher T_1 , k : - faster (less seeds), - less sensitive (some hits will be missed)
 - Lower T_1 , k : - slower (more seeds), - more sensitive (less hits will be missed)
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Exercise 3 - Programming assignment

For the programming tasks, please follow the instructions given in GitHub Classroom under the following link.

<https://classroom.github.com/a/nxAqfoYx>
