

Assignment 1 Hamming distance of two DNA sequences

- a.
- b. The Hamming distance HD for two strings s and t of the same length, say len_s , is defined as the number of character positions in which they differ. So we can compute HD by a loop in which we compare the character at each position i of the two strings, and increment HD by 1 for each position where the characters are not the same. HD must be initialized with the value 0. The corresponding Matlab code is:

```
HD=0;
for i=1:len_s
    if (s(i)~=t(i)) HD=HD+1;
    end
end
```

After reading the input file first a check is done whether the input strings are of the same length. The complete code of the file `hamming1.m` is given in Appendix A on page 3.

- c. Running the program `hamming1.m` in Matlab gives a value $\text{HD}=22$. This is indeed the correct value, as can be seen by inspecting the two strings in the input file `input.txt`.
- d. To compute the positions where '|' symbols or spaces have to be inserted, we can extend the loop above. We introduce a new string, say v : whenever there is a match on position i , we put $v(i)$ equal to '|', otherwise $v(i)$ is put equal to a space. In Matlab code:

```
for i=1:len_s
    if (s(i)~=t(i)) HD=HD+1; v(i)=' ';
    else v(i)='|';
    end
end
```

To display the strings s , v , t below one another, we can use Matlab's `print` function for each string. An alternative way is the following. We define a matrix A with 3 rows, each of length len_s , where the first row of A equals the string s , the second row equals the string v , and the third row equals the string t . Then we can use the `disp()` function of Matlab to display the matrix A , which will show the desired alignment.

The complete code of the file `hamming2.m` is given in Appendix B on page 4.

- e. Running the program `hamming2.m` gives the following output (copied from the command window of Matlab):

```
GGTCCAATGGGATTATGGCCTCTCTATATTATCCA
|   | |   ||   ||   ||   ||
GTCACCAACTTCTTTATATCTGGCTAGCTTAGATT
```

which indeed is correct.

- f. To print the alignment, which is defined by the $3 \times \text{len_s}$ matrix A , to a file `output`, we use the `fprintf` command with the output file as the first argument. The `%s` parameter indicates we are printing characters. In pseudocode:

```
for i=1:3
    for j=1:len_s
        fprintf(output, '%s', A(i, j));
    end
    fprintf(output, '\n');
end
```

The complete code of the file `hamming3.m` is given in Appendix C on page 5.

Running the extended program `hamming3.m` gives the output file `hamming3-output.txt` which is included in Appendix D on page 7.

It indeed contains the correct information.

The requested files `hamming1.m`, `hamming2.m`, `hamming3.m`, and `hamming3-output.txt` are contained in the subdirectory `results` of this directory.

A Appendix: Matlab code of hamming1.m

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Introduction to Scientific Computing - WBCS14003                      %
%                                                                       %
% Compute the Hamming distance for 2 sequences in Matlab              %
%                                                                       %
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

clear all; % remove items from the workspace

in=fopen('input.txt'); % open file
s=fgetl(in); % read line 1 of the input
t=fgetl(in); % read line 2 of the input
fclose(in); % close file
len_s=length(s); % length string s
len_t=length(t); % length string t

% Here comes your code
% First test whether the input strings are of the same length. If not, stop.
if (len_s ~= len_t)
    disp('lengths of input strings are not equal');
    return;
end

HD=0; % initialize the Hamming distance to zero

% Increase HD for each mismatch

for i=1:len_s
    if (s(i)~=t(i)) HD=HD+1;
end
end

% Print the Hamming distance on the screen:

fprintf('Hamming distance=%d\n',HD);
```

B Appendix: Matlab code of hamming2.m

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Introduction to Scientific Computing - WBCS14003                                %
%                                                                                   %
% Compute the Hamming distance for 2 sequences in Matlab                         %
%                                                                                   %
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

clear all;                                % remove items from the workspace

in=fopen('input.txt');                    % open file
s=fgetl(in);                              % read line 1 of the input
t=fgetl(in);                              % read line 2 of the input
fclose(in);                              % close file
len_s=length(s);                          % length string s
len_t=length(t);                          % length string t

% Here comes your code
if (len_s ~= len_t)
    disp('lengths of input strings are not equal');
    return;
end

% The result should be a real number HD equal to the Hamming distance of s and t

HD=0; % initialize the Hamming distance to zero

% Put string s into first row and t into third row of matrix A
A(1,:)=s;
A(3,:)=t;
% Put '|' symbol at all positions of the second row of matrix A
% where there is a match between the letters in string s and string t
% Also, increase HD for each mismatch

for i=1:len_s
    if (s(i)~=t(i)) HD=HD+1; A(2,i)=' ';
    else A(2,i)='|';
end
end

% Print the Hamming distance on the screen:

fprintf('Hamming distance=%d\n',HD);

% Print the alignment on the screen:

fprintf('\nAlignment:\n\n');
disp(A);
```

C Appendix: Matlab code of hamming3.m

```
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%
% Introduction to Scientific Computing - WBCS14003
%
% Compute the Hamming distance for 2 sequences in Matlab
%
%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%%

clear all;                                % remove items from the workspace

in=fopen('input.txt');                    % open file
s=fgetl(in);                              % read line 1 of the input
t=fgetl(in);                              % read line 2 of the input
fclose(in);                              % close file
len_s=length(s);                         % length string s
len_t=length(t);                         % length string t

% Here comes your code
if (len_s ~= len_t)
    disp('lengths of input strings are not equal');
    return;
end

% The result should be a real number HD equal to the Hamming distance of s and t

HD=0; % initialize the Hamming distance to zero

% Put string s into first row and t into third row of matrix A
A(1,:)=s;
A(3,:)=t;
% Put '|' symbol at all positions of the second row of matrix A
% where there is a match between the letters in string s and string t
% Also, increase HD for each mismatch

for i=1:len_s
    if (s(i)~=t(i)) HD=HD+1; A(2,i)=' ';
    else A(2,i)='|';
end
end

% Print the Hamming distance on the screen:

fprintf('Hamming distance=%d\n',HD);

% Print the alignment on the screen:

fprintf('\nAlignment:\n\n');
disp(A);

% Print the alignment to a file:
output=fopen('hamming3-output.txt', 'w'); % open file
```

```

fprintf(output,'Name: Anonymous\n');           % enter your name(s)
fprintf(output,'IBC, Practical Example\n');

fprintf(output,'Hamming distance=%d\n',HD);
fprintf(output,'\nAlignment:\n\n');
% Here comes the code for printing the alignment
% That is, the rows of matrix A below one another
for i=1:3
    for j=1:len_s
        fprintf(output,'%s',A(i,j));
    end
    fprintf(output,'\n');
end
fclose(output);                               % close file

```

D Appendix: output file hamming3-output.txt

Name: Anonymous
ISC, Practical Example
Hamming distance=22

Alignment:

```
GGTCCAATGGGATTATGGCCTCTCTATATTATCCA
|   | |   ||   ||   ||   ||
GTCACCAACTTCTTTATATCTGGCTAGCTTAGATT
```