

## Practical 9

### Questions

21<sup>st</sup> March 2022

1. Identify the pair of sequences which are close to each other using Hamming and Euclidean distance methods.

(i) AMENLNMDLLYMAAAVMMGLAAIGAAIGIGILGGKFLEGAARQPDLIPLLRTQFFIVMGLVDAIPMIAVG  
LGLYVMFAVA

(ii) AADVSAAVGATGQSGMTYRLGLSWDWDKSWWQTSTGRLTGYWDAGYTYWEGGDEGAGKHSLSFAP  
VFVYEFAGDSIKPFIEAGIGVAAFSGTRVGDQNLGSSLNFEDRIGAGLKFANGQSVGVRAIHYSNAGLKQPN  
DGIESYSLFYKIPI

(iii) MALLPAAPGAPARATPTRWPVGCENRPWTKWSYDEALDGIKAAGYAWTGLLTASKPSLHHATATPEY  
LAALKQKSRHAA

2. Get the non-redundant sequences of beta barrel membrane proteins with sequence identities of less than 40%, 50%, 75% and 90% using CD-HIT

3. Get the non-redundant sequences of the same type of proteins with sequence identities of less than 20%, 30%, 40% and 50% using PISCES.

4. Compare the results obtained with the cut-offs 40% and 50%.

5. Extract the data with the cut-off of 50% from Uniprot and compare with CD-HIT and PISCES.

**Deadline: 27<sup>th</sup> March 2022**