

Project 4

GOR III

Step 1 - parser

- parse file CATH_info.txt
 - first four characters correspond to the file name
 - the fifth character to the chain that should be used
- parse files in dssp directory
 - only take those chains indicated in the fifth character in CATH_info.txt
 - check columns 3 to 5 to recuperate sequence, AA and corresponding class
 - regroup class information
 - H, G, I -> H
 - E, B -> E
 - T -> T
 - C, S, “ “ -> C

Step 2 - GOR III

- general formula

$$I(\Delta S; R) = I(S; R) - I(n-S; R) = \log(f_{S,R}/f_{n-S,R}) + \log(f_{n-S}/f_S) \quad (3)$$

- GOR III adaptation

$$I(\Delta S_j; R_1, \dots, R_n) \approx I(\Delta S_j; R_j) + \sum_{m, m \neq 0} I(\Delta S_j; R_{j+m} | R_j) \quad (8)$$

$$I(\Delta S_j; R_{j+m} | R_j) = \log(f_{S_j, R_{j+m}, R_j} / f_{n-S_j, R_{j+m}, R_j}) + \log(f_{n-S_j, R_j} / f_{S_j, R_j}) \quad (9)$$

- count the following
 - frequency of the structure (f_S)
 - frequency of the pair (structure, AA: $f_{\{S, R\}}$)
 - frequency of the triplet (structure, AA neighbor, AA: $f_{\{S, R_m, R\}}$)
 - neighborhood of 8 AA to the left and 8 aa to the right
- use the frequencies to compute predictions: that conformation S with the highest value in equation (8) will be the predicted conformation

Step 3 - Quality of predictions

- compute Q3 and MCC
 - **Q3**: number of correctly predicted residues/total number of residues
 - **MCC**

$$\text{MCC} = \frac{TP \times TN - FP \times FN}{\sqrt{(TP + FP)(TP + FN)(TN + FP)(TN + FN)}}$$

- visualize your prediction quality using ROC curve(s)