

Using a hidden Markov model for prediction of transmembrane helices

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Introduction

- Given: Dataset consisting of 160 membrane proteins annotated with their transmembrane helices
- Assignment: Train a 3 state and 4 state model by training parts 0-8 dataset using "training-by-counting"
- Assignment: Use 3 and 4 state model to make a 10-fold-experiment, training-by-counting and viterbi decoding for prediction.

Train the 3-state model

```
# Go through every sequence, matching it up with the annotation, counting the states
for index, seq in enumerate(sequences):
    sequence_index = [observable_to_index[observation] for observation in seq]
    annotation = [states_to_index[c] for c in sequence_annotations[index]]
    for j, amino_acid in enumerate(sequence_index):
        if j == 0:
            # First one, so count in pi_table
            pi_table[annotation[j]] += 1
            emissions_table[annotation[j]][amino_acid] += 1
        if j > 0:
            # Can get a transition from past value to current one
            transitions_table[annotation[j-1]][annotation[j]] += 1
```

Train 4-state model

- Preprocessing annotation
 - from "o" to "M" = "m"
 - from "i" to "M" = "M"
- Go through every sequence, matching it up with the annotation and counting the states (just as 3 state model)

10-fold experiment

```
for test in range(0, 10):  
    observables = []  
    sequences = []  
    sequence_annotations = []  
    sequence_names = []  
    spot = 1  
    for data_file_num in range(0, 10):  
        if data_file_num != test:
```

Training-by-counting

- 3-state model, for parts 0-8:

Start Probabilities:

```
[ 0.5170068  0.00680272  0.47619048]
```

Transition Probabilities:

```
[[ 9.80093452e-01  1.98425398e-02  6.40081930e-05]
 [ 2.33909632e-02  9.54089786e-01  2.25192503e-02]
 [ 3.56531660e-05  1.14803195e-02  9.88484027e-01]]
```

Emission Probabilities:

```
[[ 0.08056209  0.01214472  0.06727284  0.04946907  0.06886247  0.03446303
  0.04304699  0.02339925  0.07013416  0.02797736  0.08266039  0.04063076
  0.04044001  0.0527119  0.07146945  0.08329624  0.05404718  0.01316208
  0.05760794  0.02664208]
 [ 0.10984546  0.01879126  0.00711021  0.00717297  0.07915548  0.08858739
  0.11782631  0.00856127  0.00551404  0.04062976  0.16513096  0.01646956
  0.00921425  0.02938402  0.05470507  0.0058768  0.05122252  0.02836828
  0.11514184  0.04099253]
 [ 0.06564256  0.02290022  0.0600953  0.05543702  0.07108314  0.04160444
  0.04651163  0.02268686  0.04644051  0.02410924  0.08847166  0.05056539
  0.04256454  0.05700164  0.07079866  0.04988977  0.06294005  0.0195932
  0.06382903  0.03783515]]
```

Training-by-counting

- 4-state model, for parts 0-8:

Start Probabilities:

```
[ 0.51351351  0.00675676  0.00675676  0.47297297]
```

Transition Probabilities:

```
[[ 9.80030722e-01  1.98412698e-02  6.40040963e-05  6.40040963e-05]
 [ 1.49543891e-04  9.53342306e-01  1.49543891e-04  4.63586063e-02]
 [ 4.54545455e-02  1.41163185e-04  9.54263128e-01  1.41163185e-04]
 [ 3.56518949e-05  3.56518949e-05  1.14799102e-02  9.88448786e-01]]
```

Emission Probabilities:

```
[[ 0.08056209  0.01214472  0.06727284  0.04946907  0.06886247  0.03446303
  0.04304699  0.02339925  0.07013416  0.02797736  0.08266039  0.04063076
  0.04044001  0.0527119  0.07146945  0.08329624  0.05404718  0.01316208
  0.05760794  0.02664208]
 [ 0.11129345  0.015963  0.00775772  0.0068626  0.0787707  0.08846785
  0.114128  0.00671341  0.0056691  0.0399821  0.16843205  0.01670894
  0.00835447  0.0331195  0.0547516  0.00760853  0.05504998  0.02924064
  0.11487394  0.03625242]
 [ 0.10830986  0.0215493  0.00661972  0.00816901  0.07943662  0.08859155
  0.12112676  0.01042254  0.00549296  0.04126761  0.16169014  0.01633803
  0.01014085  0.02591549  0.05464789  0.0043662  0.04760563  0.02760563
  0.11521127  0.04549296]
 [ 0.06564256  0.02290022  0.0600953  0.05543702  0.07108314  0.04160444
  0.04651163  0.02268686  0.04644051  0.02410924  0.08847166  0.05056539
  0.04256454  0.05700164  0.07079866  0.04988977  0.06294005  0.0195932
  0.06382903  0.03783515]]
```

3-State Viterbi Results

- Over all 10 folds:

Variance: 0.00442402539894

Mean: 0.676804974853

- Individual ACs:

0.630622152503856

0.7588685678999825

0.6512626924662515

0.5701947755526338

0.7101011545096947

0.713966925168569

0.7002301736031689

0.6545562990011426

0.5903359974642108

0.7879110103589546

4-State Viterbi Results

- Over all 10 folds:

Variance: 0.00468726047895

Mean: 0.680697300092

- Individual ACs:

0.5825665066458083

0.7400915818648046

0.643551590962399

0.6066165791228548

0.7359067133402921

0.7177319941203848

0.7404435898416963

0.6368863895360404

0.6119622374686893

0.7912158180171618