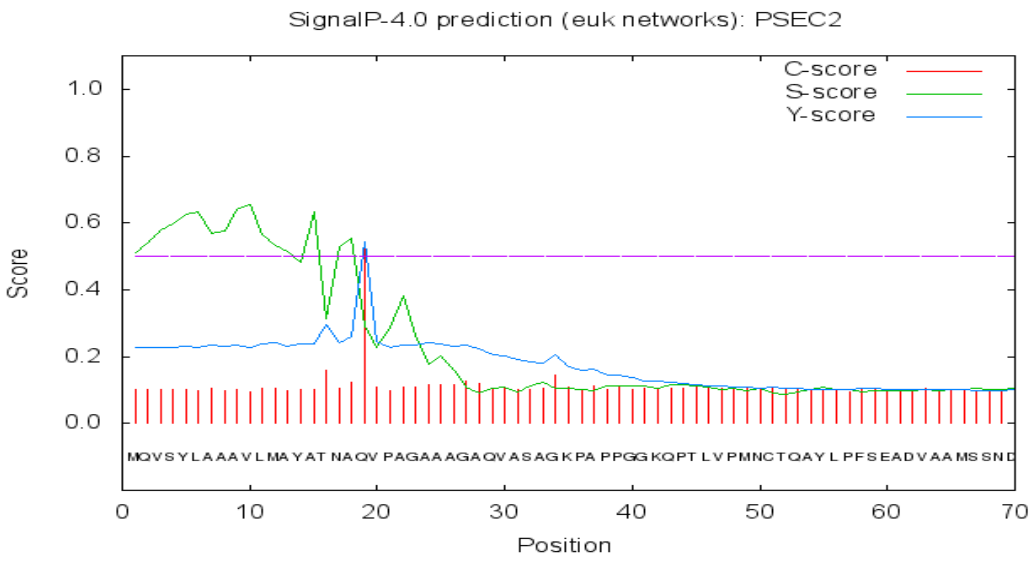
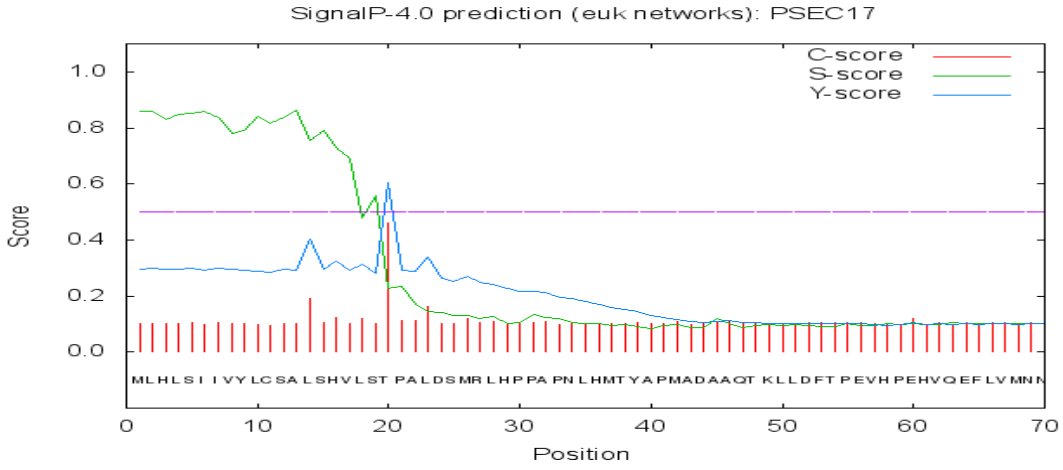
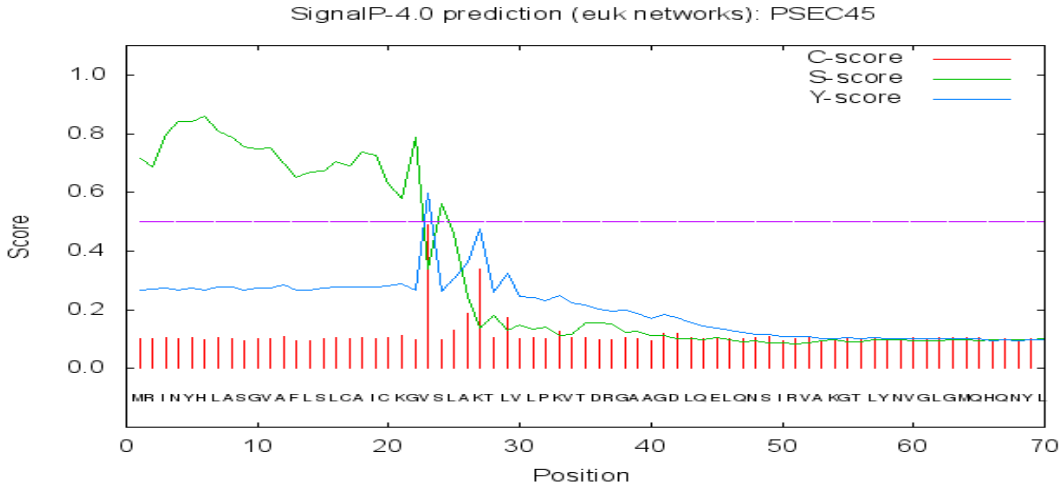
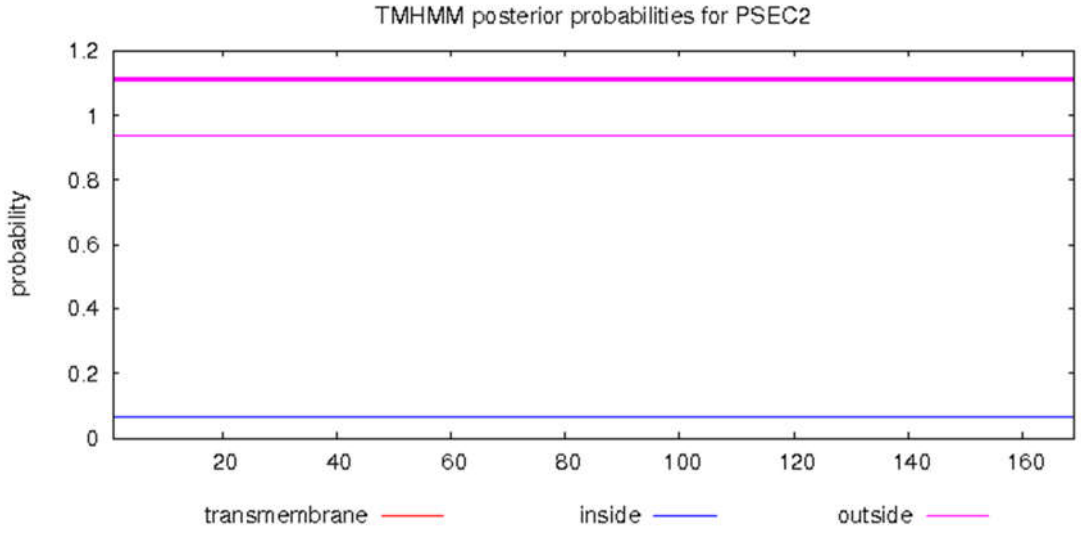
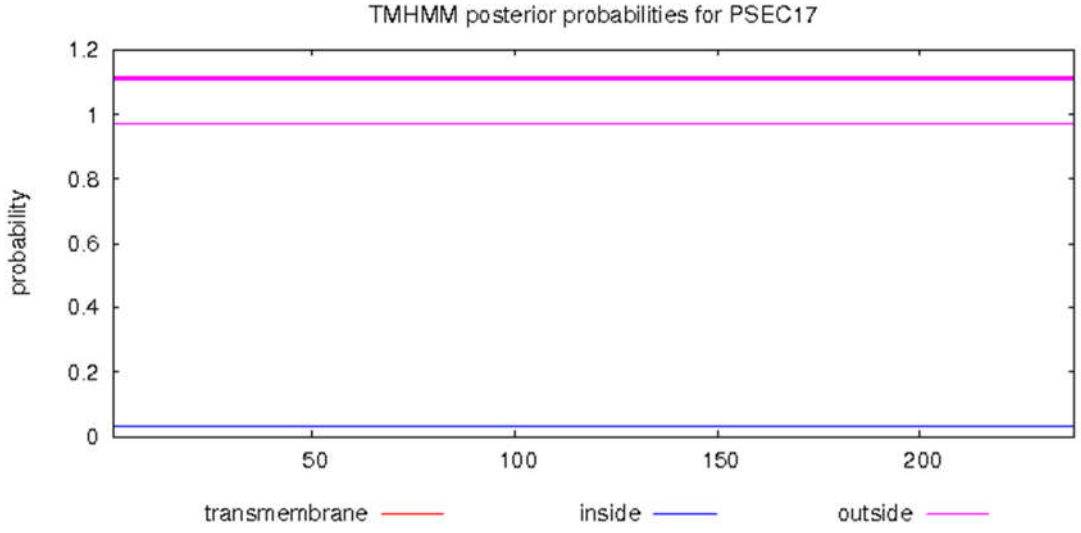
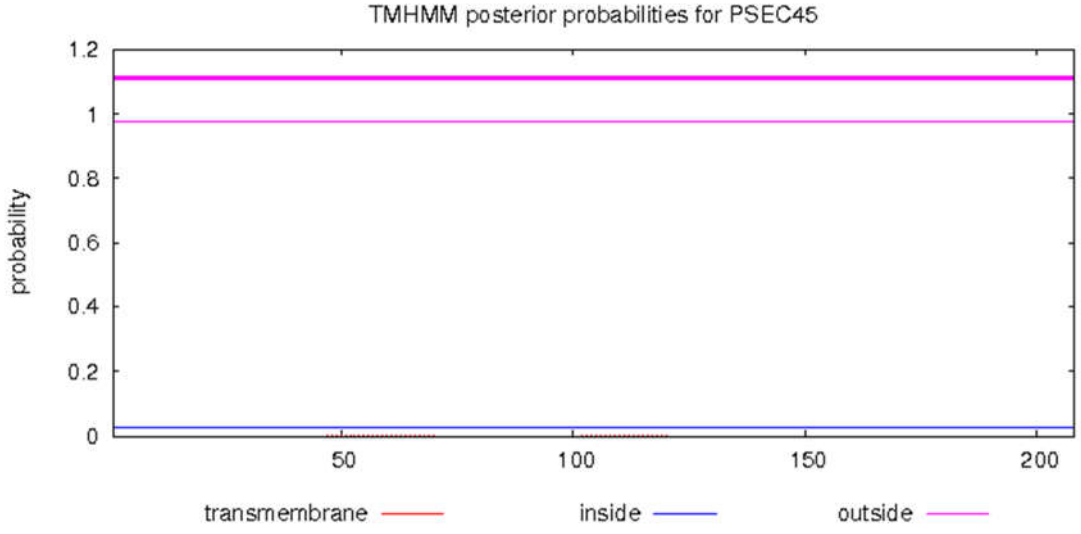

SignalP-4.0 prediction (euk networks): PSEC2
C-score
S-score
Y-score
Score
Position
MQVSYLAAAVLMAYATNAQVPAGAAAGAQVASAGKPAPPGGKQPTLVPMNCTQAYLPFSEADVAAMSSND


SignalP-4.0 prediction (euk networks): PSEC17
C-score
S-score
Y-score
Score
Position
MLHLSIIVYLCSALSHVLSTPALDSMRLHPPAPNLHMTYAPMADAAQTKLLDFTPEVHPEHVQEFLVMNN


SignalP-4.0 prediction (euk networks): PSEC45
C-score
S-score
Y-score
Score
Position
MRINYHLASGVAFLSLCAICKGVSLAKTLVLPKVTDRGAAGDLQELQNSIRVAKGTLYNVGLGMQHQNYL


TMHMM posterior probabilities for PSEC2
probability
1.2
1
0.8
0.6
0.4
0.2
0
20
40
60
80
100
120
140
160
transmembrane
inside
outside


TMHMM posterior probabilities for PSEC17
probability
1.2
1
0.8
0.6
0.4
0.2
0
50
100
150
200
transmembrane
inside
outside


TMHMM posterior probabilities for PSEC45
probability
1.2
1
0.8
0.6
0.4
0.2
0
50
100
150
200
transmembrane
inside
outside