蛋白质跨膜结构预测分析

# WEBSEQUENCE Length: 200

# WEBSEQUENCE Number of predicted TMHs: 0

# WEBSEQUENCE Exp number of AAs in TMHs: 6.18768

# WEBSEQUENCE Exp number, first 60 AAs: 0

# WEBSEQUENCE Total prob of N-in: 0.10994

WEBSEQUENCE TMHMM2.0 outside 1 200