

Dmel7883\_9-170 ENNKVFVLIILLYCVLVSILKLCTAQPDSSVAATDNDITHLGDDCQVTPVIVHLQYPGCVPKPIPSFACVGRCASYSIQVSG 80

Consensus



Dmel7883\_9-170 SKIWQMERSCMCCQESGEREAAVSLFCPKVKPGERKKVLTAKAPLECMCRPCTSI EESGII PQEIAGYSDEGPLNNHFR 160

Consensus



Dmel7883\_9-170 RI 162

Consensus

