

Dmel8553_1-288 MRPLSGLIALALLLLLLLTAPSSAADTETESSGSPLTPGAEEP RR VVKRAPTSSFI GMRGKKDEEHDTSEGNWLGSGPD P 80

Consensus



Dmel8553_1-288 LDYADEEADSSYAENGRR LKKAPLAFVGLRGKKFIPINNRLSDVLQSLEEEERLRDSSLQDFFDRVAGR DGS AVGKRAPT G 160

Consensus



Dmel8553_1-288 FTGMRGKR PALLAGDDDDAEADEATELQQKRAPVNSFVGMRGKKDVS HQHYKRAALSDSYDLRGKQQRFA DFN SKFVAVR G 240

Consensus



Dmel8553_1-288 KKSDLEGNGVGIGDDHEQALVHPWLYLWGEKRAPNGFLGMRGKR PALF 288

Consensus

