

Dmel2661\_1-102 MARYTYLVALLAICCWGYCGAMAMPDEERYVRKEYNRDLLDWFNNVGVGQFSPGQVATLCRYPLILENSLGPSPVPIR 80

10 20 30 40 50 60 70 80

Consensus



Dmel2661\_1-102 KRNSELINSLSLPKNMNDAGK 102

90 100

Consensus

