

Dpul10065\_1-145 MKGKGAF LMVLAGWGLIGLMILT TAVEEAAPH PADY TSSSVNNQRDFRSRRGF KTVGLATARGFGKRAPSL SNFNSFQDAA 80

Consensus



Dpul10065\_1-145 EQMMQQQEENPN SDPDVFPVDWLVN YLQNKPDVIRYMVEHLLDHNGDGQVTSQEMMTSLQQQRED 145

Consensus

