

Dmel11489\_1-207 MAHTKTRRTYGF L MVLL I LGSACGNLVA S GSAGSP PSNE PGGGGLSEQVVL DQLSESDLYGN NKRAWQS LQSSW GKRSSS 80

Consensus



Dmel11489\_1-207 GDVSDPD IYMTGHFVPLV I TDGTNT I DWDTFERL A SGQSA QQQQQQP LQQQS QSGEDFDD LAGEPDVEK RAWKSMNV AWG 160

Consensus



Dmel11489\_1-207 KRRQAQGWNKFRGAWGKREPTWNN LKGMW GKR DQWQKLHGGW GKR SQ 207

Consensus

