

Tmol15165_1-82 MFDRKFFVFAVFLVVVFATLALLETSTRHTVEARYLPTRSNNGDRVDKLRRELLKDLLQSEIEKEEYQADAPQRWHPENKLFYKREA 80
Tmol14259_1-57 MFARKFFFFTACLVLFVSLALETSHTVHARYLPTRSNNGDRVDKLRRELLRDILLEKELD - D - - - - - - - - - - - - - - - - - - - - 57
Tmol3528_1-74 MFARKFFFSAACLVLFVTLALLETSTRHTVHARYLPTRSNNGDRVDKLRRELLRDMLEKELD - DEYQGYVISRMHPENKLS - - - - - - - - - - 74

Consensus



Tmol15165_1-82 PA 82
Tmol14259_1-57 - - 57
Tmol3528_1-74 - - 74

Consensus

