

Tmol18276\_1-95

80

10 20 30 40 50 60 70

MQGYSAAMKCLCAVTCIMIVVATVTS AAPSYADYDNNIRDLWEMLLQKDALEDKFGGHQMVRKSSRS PSLRLRFGRRS DSS

Consensus

MQGYSAAMKCLCAVTCIMIVVATVTS AAPSYADYDNNIRDLWEMLLQKDALEDKFGGHQMVRKSSRS PSLRLRFGRRS DSS

Tmol18276\_1-95

95

90

MSPEAAFMMAQAVDH

Consensus

MSPEAAFMMAQAVDH