

Tcas2233_1-136 MKTFLIYSACVVLFCIANCQGEPEPKRNKLASVYALTTPSLRVGRRSEGTDVKRRIGKMVSFPRIGRSESNWVPDDNSYG 80

10 20 30 40 50 60 70 80

Consensus



Tcas2233_1-136 AQRPGANS GG MWFGPRLGRVQKRSENF T PWAYI I L NGEA P I I R E V H Y S P R L G R E S E 136

90 100 110 120 130

Consensus

