

Tcas13749_17-149 VDDRCDNEICQNNFDDLIKRKVKDNEDMTYQSTTRLSPYYHPSRPMPCSCGIEFRVLDLGHQYYPRYLHSGVCKSELCCG - 79
Tcas13253_117-211 - - - - - QQWHCKSKIET - - DLGPDYFPRYLKSVVCLSENCCWF 35

10 20 30 40 50 60 70 80

Consensus



Tcas13749_17-149 GPYRCIERHYKVRVLKQKD - - - - - PRNPEIRPSMALPDTLKGTLWSETITVTVACECSV 133
Tcas13253_117-211 GLYKCKPRSFVNLRLRRKKGFCAANGGKNNHKVGVAGLPNDLKELVWWEERAVNFCDCR 95

90 100 110 120 130 140

Consensus

