

Tcas11245\_5-112 NGISLLMLLVAGMILFQA TTTYAAPHSPSYPGYYSPISMEGQNPEYLLQT IARLRQALISDDDDLENSKRGLDLGLGRGFS 80

Consensus



Tcas11245\_5-112 GSQAAKHLMGLAAANFAGGPGRRRRSE E 108

Consensus

