

MAAVVLENGVLSRKLSDFGQETS... 120
MAAVVLENGVLSRKLSDFGQETS... 120
MSTAVLENPGLGRKLSDFGQETS... 120
MSALVLESRALGRKLSDFGQETS... 119
... 0
... 0
... 0
... 0
... 0

FPRTIQELDRFANQILSYGAEL... 240
FPRTIQELDRFANQILSYGAEL... 240
FPRTIQELDRFANQILSYGAEL... 240
FPRTIQELDNFANQVLSYGAEL... 239
... 93
... 109
... 84
... 76
... 90

RLRPVAGLLSSRD... 358
RLRPVAGLLSSRD... 358
RLRPVAGLLSSRD... 358
RLRPVAGLLSSRD... 357
KIVAVPGLIPDDV... 212
QIVAVPGLVPDEV... 228
TVVAVPGLVPDDV... 203
RVARVPALIPFQT... 194
EIIAVPGLIPAAP... 210

SD.KPKLLPLELEKTACQEY... 453
SD.KPKLLPLELEKTACQEY... 453
SE.KPKLLPLELEKTAIQNY... 452
SD.KPKLLPLELEKTAVQEY... 451
DSASP... 297
DSASP... 313
DDPSP... 294
SD.EPLHQAFNPLEAMRT... 262
EGKDAH... 275

[X] non conserved
[X] ≥ 50% conserved