Multiple sequence alignment of species: Isca, Vdes, Dpul, Apis, Tcas, Phum, Aaeg, Dmel, Smar, Lmig, Bmor, Znev, Tmol8433_9-221:., Tmol19835_8-182:.

Consensus (≈1–80):
MSPVESSRHAGRRPVVATYGESGXMMRXAHFXLAXVLGALLXLXVLXSXXGDPXSQTXAVELHQQXSGXXSSXXSXQSXXX

Consensus (≈81–160):
QXQVDEDXEDKRDWRXLXXGWXKRXWQDDLHGWGKRXGWXNLXQGGWGKRRSXPGWEDPQSXAWGKRXXQDLXGGWGKRA

Consensus (≈161–240):
XDTWQDLXQSGWGKRDXXXXXLXKGWQDLQXFWXKEDWYDLRGGAXWMGMRGKRXWXXLXNGGWGKRGDAXYYXXDXXXX

Consensus (≈241–320):
XXXXXXXXXXXXXXXXDEGDVWGXRDWXDXXNDWXKRAWNXFQGXWGKRXPXXSKTNSGPAIEGMGNNDLLLLISGTGDQL

Consensus (≈321–400):
YQQREDDQAKANVEDAGNEESVDKRGWNNFXGSWGKRDPXXPDXEXXXXXXXXVXXXXXXEXIGAPXKRAWNNLKGXW

Consensus (≈401–470):
GKREIPXXXXXXXXXXXXXKRXXGWNNLKGLWGKRADTNWNRLSAAWGKRSIGGETGIKEXXXXXTSXEKIIIII