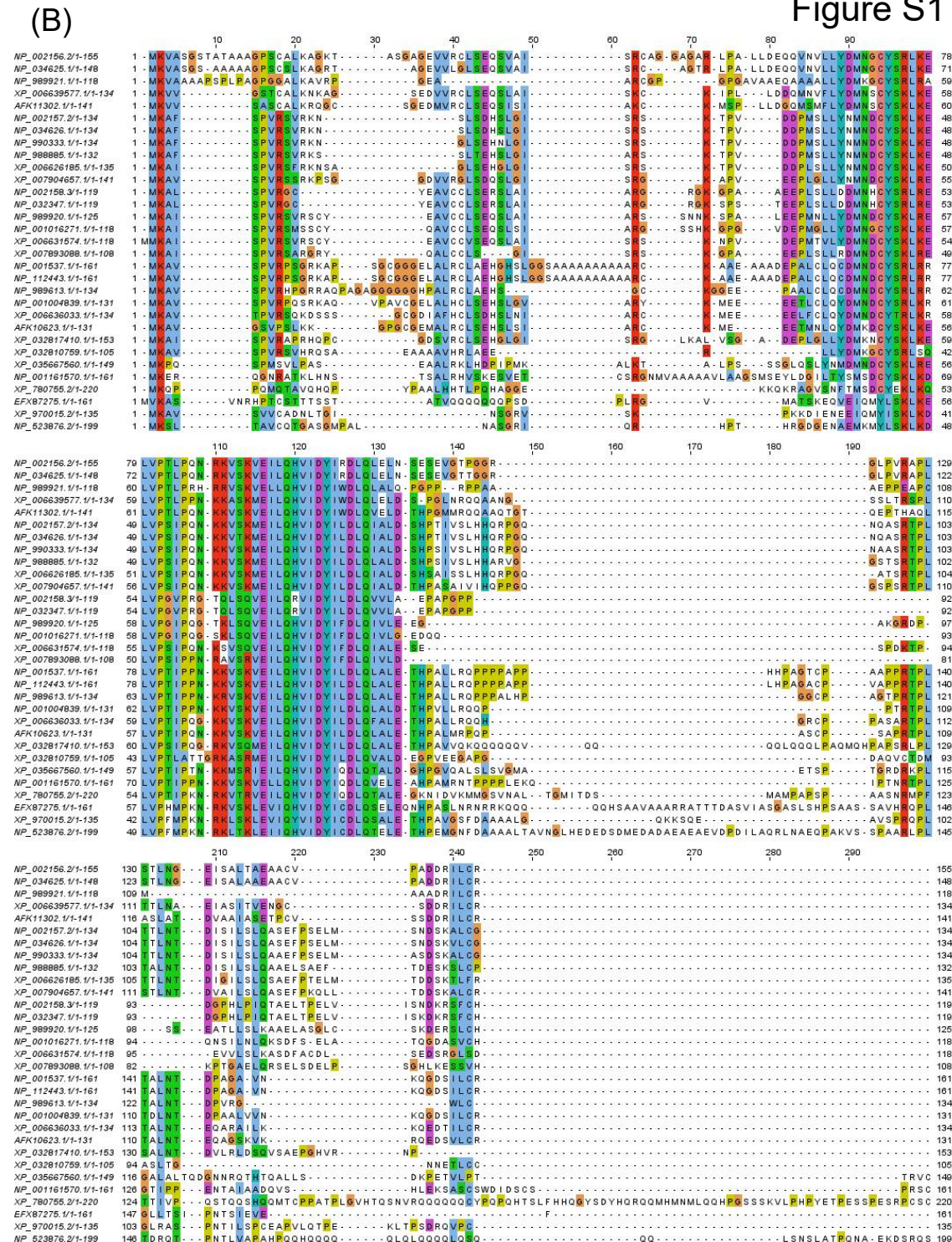(A)

Figure S1. Multiple sequence alignments of E proteins (A) and Id proteins (B). The alignments were visualized using Jalview (version 2.11.1.4) [1].

1. Waterhouse, A. M.; Procter, J. B.; Martin, D. M.; Clamp, M.; Barton, G. J. Jalview Version 2 - a multiple sequence alignment editor and analysis workbench. Bioinformatics **2009**, 25(9), p. 1189-1191.