

AlpA	1	-----MNN--RSAVRILRLPAVIQKTGMARATIYDNLNPKSPRYDATEFP-KKRMIGVKS VGWIEAEIDEWLSQRCKLI---	70
Hef	1	MHPLFRAHQLSLRLPLCFFILRPPFISVTGGYMTSYQLRLRQVEQKTGLKRSQIYLYMK-----EGAFP-RSIKIGPASVAWLESEIDEWINKKLSDR---	61
VefB	1	-----MQLTNLRGVNMPENNIRLIRFREVLTMTGLSRSSLYRFIE-----ENQFP-PQVQTGGRAVAWVEGEVQEWIAQRITNRRMD	76
VefA	1	-----MGQQDIGEHPMRFLKLKEVMEKTALSRSAIYRKMN-----DGEFP-QSVSLGERAIAWVESEVDEWMDFCLKQR---	68
XisSXT	1	-----MQMTAVAHQVTPFLTSTYEVMARYHISYTTLWRRRIK-----DGSLEFPQPRINRNTRNKLWHIEDLEEYEKK--ED---	66
consensus	1	i m m vrilrlrevi ktglsrstiyrl dg fP i lg r vaWie eidEwl rl r	