

[illegible][illegible][illegible]

NTDB id 361	SMU RS01310 WP 002262737.1	YGTVDDEVFYNPQHPPYTWGLLNSMPTTNTAS	GSLSHSIPGTPPDLLQPPQGDAFAPRNEFALDIDLKEPPFFKVS	DSHYAA	316
NTDB id 977282	MGAS37850 RS08210 WP 010921893.1	FGTVEEIFYDPRHPYTWSLLSSLPQLADESGEL	YAIAGTPPSPLYSPIIGDAFALRSEYAMVLDFEKAPPA	INVSETHWAK	320
NTDB id 478	HSIS4 RS06985 WP 021143811.1	YATVEEIFYEPCHPYTWSLLSSLPQLADDKG	ELFSIPGTPPSLYTTPVVGDAFALRSDYAMQIDFEEKAP	QFQVSDTHWAK	320
NTDB id 327	STU RS16115 WP 011226304.1	YATVEEIFYEPCHPYTWSLLSSLPQLADDNGK	LFSIPGTPPSLYTTPVVGDAFALRSDYALQIDFEEKAP	QFQVSDTHWAK	320
NTDB id 295	STER RS06915 WP 011226304.1	YATVEEIFYEPCHPYTWSLLSSLPQLADDNGK	LFSIPGTPPSLYTTPVVGDAFALRSDYALQIDFEEKAP	QFQVSDTHWAK	320
consensus		*** ** ** ** **	*** ** ** ** **	*** ** ** ** **	*** ** ** **

NTDB id 361	SMU RS01310 WP 002262737.1	TWLLD D RAPK I TPPERILQ L RWEKWKSLSQGEEKHND.....	350
NTDB id 977282	MGAS37850 RS08210 WP 010921893.1	TWLLHP E A P KVQKP E VIVQLDHQKILRKMSQQEEGV....	356
NTDB id 478	HSIS54 RS06985 WP 021143811.1	TWLLHEDAPKV E KPAVIQN L HEKILRANMGMFAHLGDEEEEGNA	361
NTDB id 327	STU RS16115 WP 011226304.1	TWLLHEDAPKV D KPAVIQN L HEKILANMGMFAHLGDEEEEGNA	361
NTDB id 295	STER RS06915 WP 011226304.1	TWLLHEDAPKV D KPAVIQN L HEKILANMGMFAHLGDEEEEGNA	361
consensus		***** ** *** ***** ***** ***** *****	

 non conserved
 similar
 $\geq 50\%$ conserved