

NTDB id 146	SP RS08570 WP 000614538.1	.IKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	.AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396
NTDB id 216	SPD RS08205 WP 000614552.1	.IKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	.AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396
NTDB id 182	SPR RS07820 WP 000614552.1	.IKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	.AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396
NTDB id 257	KZH43 RS07655 WP 220041236.1	.IKNPSQAVTEETYQPQAPKKHRFKMRYLILLASLVLV.	.AASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGE	396
NTDB id 467	HSISS4 RS06915 WP 021143821.1	PTPAPSKQPRKKTTPAKKKKKKKNNFFSTLLKVVFLGLVFI	LGIIIFAYLVFTNPDNVQVPNVVVGQELSTAQTKLEGAGFKVGE	396
NTDB id 1126901	FGK94 RS08340 WP 043030078.1	ARLEQTSQMDTLQKPQKKKKNGRFLGTLKILFSFFVVGVALFAYLVLT	PTTVKVPDVTGSSSLNAAKQVLEEAGLKVCK	398
NTDB id 384	SMU RS02325 WP 002263039.1	DKVVDHKSCKPKTK.PQPKPKKKRRFLARFFKIFFILVAIGIAVL	LYLVLTKPTSTVSVDPVSGDKLSTAKMIIRKSSGLKVGD	396
consensus		*****	*****	*****

[illegible][illegible]

NTDB id 146	SP RS08570 WP 000614538.1	SESEPGTIMKQSPGAGTGYDVKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTPAGS	A	E	624	
NTDB id 216	SPD RS08205 WP 000614552.1	SESEPGTIMKQSPGAGTGYDVKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTPAGS	V	E	624	
NTDB id 182	SPR RS07820 WP 000614552.1	SESEPGTIMKQSPGAGTGYDVKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTPAGS	V	E	624	
NTDB id 257	KZH43 RS07655 WP 220041236.1	SESEPGTIMKQSPGAGTGYDVKPTQIVLTVAKKVTSVAMPSYIGSSLEFTKNNLIQIVGIKEANIEVVEVTTPAGS	V	E	624	
NTDB id 467	HSISS4 RS06915 WP 021143821.1	IPSSSTE....ILYQDPQAGTSVDGT	VILYVS	SVATASSSLQSS.....SS	593	
NTDB id 1126901	FGK94 RS08340 WP 043030078.1	SPSSSAVVSGQTPIYYGTQISLS	SEKGD	ISLYLYPEET	RASSSSS.....SS	601
NTDB id 384	SMU RS02325 WP 002263039.1	SPSSSLVVGQSPYYGNTVSLSS	SNDD	IILYVSTSGGS	SHSGSS.....S	591
consensus		**!*!!!!!!!!!!!!!!!!!!!!!!*!*****!*****!*****!*****!*****!				*

logo

NTDB id 146 SP RS08570 WP 000614538.1	GMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP..	659
NTDB id 216 SPD RS08205 WP 000614552.1	GMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP..	659
NTDB id 182 SPR RS07820 WP 000614552.1	GMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP..	659
NTDB id 257 KZH43 RS07655 WP 220041236.1	GMVVEQSPRAGEKVDLNKTRVKISIIYKPKTTSATP..	659
NTDB id 467 HSISS4 RS06915 WP 021143821.1	STTHSS..STSSSTDSTTSSTE....TSTEATHTELQ	624
NTDB id 1126901 FGK94 RS08340 WP 043030078.1	DTSAPN.PSAGDTTSSSSHTEL....GQNDAVTP..	631
NTDB id 384 SMU RS02325 WP 002263039.1	SSNSEGITTSSEASTDSSSSATT....TS.....	616
consensus	*****!*****	

- non conserved
- similar
- ≥ 50% conserved