

logo

NTDB id 146 SP RS08570 WP 000614538.1
NTDB id 216 SPD RS08205 WP 000614552.1
NTDB id 182 SPR RS07820 WP 000614552.1
NTDB id 257 KZH43 RS07655 WP 220041236.1
NTDB id 467 HSISS4 RS06915 WP 021143821.1
NTDB id 1136586 DQL35 RS08460 WP 111678383.1
NTDB id 384 SMU RS02325 WP 002263039.1
consensus



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NTDB id 146 SP RS08570 WP 000614538.1
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consensus



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NTDB id 146 SP RS08570 WP 000614538.1
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NTDB id 146 SP RS08570 WP 000614538.1
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consensus



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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	LGIIIFAYLVFTNPDNVQVVPNVVQGQELSTAQTKLEGAGFKVGE	396
NTDB id 1136586	DQL35 RS08460 WP 111678383.1	ARLEQTSQMDTLQKPQKKKKNGRFLGTLKILFSFFVVGVALFAYLVLT	PTTVKVPDVTGSSSLNAAKQVLEEAGLKVCK	398
NTDB id 384	SMU RS02325 WP 002263039.1	DKVVDHKSCKPKTK.PQPKPKKKRRFLARFFKIFFILVAIGIAVL	LYLVLTKPTSTVSVDPVSGDKLSTAKMIIRKSSGLKVGD	396
consensus		*****	*****	*****

NTDB id	146	SP RS08570 WP 000614538.1	EKTEASEKVEEGRIIRTPDGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK.	KVPDNLIKIEEEESNES	475
NTDB id	216	SPR RS08205 WP 000614552.1	EKTEASEKVEEGRIIRTPDGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK.	KVPDNLIKIEEEESNES	475
NTDB id	182	SPR RS07820 WP 000614552.1	EKTEASEKVEEGRIIRTPDGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK.	KVPDNLIKIEEEESNES	475
NTDB id	257	KZH43 RS07655 WP 220041236.1	EKTEASEKVEEGRIIRTPDGAGTGRKEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK.	KVPDNLIKIEEEESNES	475
NTDB id	467	HSISS4 RS06915 WP 021143821.1	VKEVEDDSVDTGKVIKTDPTAGTTTrKEGSSIDVYVSSSGSKGFALKDYKGKNYKDAIEDLTSNFGVSEDQIDIQHVEDDSA		476
NTDB id	1136586	DQL35 RS08460 WP 111678383.1	VHQIESDVTSEDDQVVKTNPQAQSAKRKCSSVDLYLSICNGKFEMENYKGGDYQDVMTSLTETYGVPKSKI KIERIVTNEY		478
NTDB id	384	SMU RS02325 WP 002263039.1	VQKIEDNVGAKGVVRTPNSAGSKRKREGSSVDIFVA.SAKGFKMEDYTGDYKDIDAINDLTNNYGVSRSIVLKEVSSDDY		475
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NTDB id 146	SP RS08570 WP 000614538.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEEEE	S	544
NTDB id 216	SPD RS08205 WP 000614552.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEEEE	S	544
NTDB id 182	SPR RS07820 WP 000614552.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEEEE	S	544
NTDB id 257	KZH43 RS07655 WP 220041236.1	EAGTVLKQSLPEGTTYDLSKA	TQIVLTVAKKATTIQLGNYIGRNSTEVISELKQKKVPENLIKIEEEE	S	544
NTDB id 467	HSISS4 RS06915 WP 021143821.1	EEGEILSSSPGKKNKSFNPKDSKAKIKFRVATPK.IVTMPDVTCLTVSTAVQTLNRKN.ISSSSIEYHDYNTGAK.LDKAK			553
NTDB id 1136586	DQL35 RS08460 WP 111678383.1	PENTVVISQSSPSAGEKFNPA.G.KSKITLSVAVG.D.TVIMPMVTEYSYADAVNTLTALGIDAARIKAYVPSSSSPTGFVQV			556
NTDB id 384	SMU RS02325 WP 002263039.1	SGCTVIGSQSPKPKGKTYHPS.S.DKKITLKV.V...KVTMPNLKNSTYEEAVSTLTAMG.ISSSR.IKAYDASDYS...SEI			546
consensus		*****!*****	*****!*****	*	

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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 1136586 DQL35 RS08460 WP 111678383.1	DTSAPN.PSAGDTTSSSSHTEL....GQNDAVTP..	631
NTDB id 384 SMU RS02325 WP 002263039.1	SSNSEGITTSSEASTDSSSSATT....TS.....	616
consensus	*****!*****	

- non conserved
- similar
- ≥ 50% conserved