

logo

NTDB id 20079 SSU05 0428 ABP89395.1
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 384 SMU RS02325 WP 002263039.1
consensus



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NTDB id 146 SP RS08570 WP 000614538.1
NTDB id 216 SPD RS08205 WP 000614552.1
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NTDB id 257 KZH43 RS07655 WP 220041236.1
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consensus



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NTDB id 20079 SSU05 0428 ABP89395.1
NTDB id 146 SP RS08570 WP 000614538.1
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consensus



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NTDB id 20079 SSU05 0428 ABP89395.1
NTDB id 146 SP RS08570 WP 000614538.1
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consensus



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NTDB id 20079 SSU05 0428 ABP89395.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	423
NTDB id 146 SP RS08570 WP 000614538.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
NTDB id 216 SPD RS08205 WP 000614552.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
NTDB id 182 SPR RS07820 WP 000614552.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
NTDB id 257 KZH43 RS07655 WP 220041236.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
NTDB id 467 HSISS4 RS06915 WP 021143821.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
NTDB id 384 SMU RS02325 WP 002263039.1	KHRFKMRYLILLASLVLVAASLIWILSRTPATIAIPDVAGQTVAEAKATLKKANFEIGEEKTEASEKVEEGRIIRTPD GAGTGR	421
consensus	*****	

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NTDB id 20079 SSU05 0428 ABP89395.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	500
NTDB id 146 SP RS08570 WP 000614538.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	504
NTDB id 216 SPD RS08205 WP 000614552.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	504
NTDB id 182 SPR RS07820 WP 000614552.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	504
NTDB id 257 KZH43 RS07655 WP 220041236.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	504
NTDB id 467 HSISS4 RS06915 WP 021143821.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	506
NTDB id 384 SMU RS02325 WP 002263039.1	KEGTKINLVSSGKQSFQISNYVGRKSSDVIAELKEK KVPDNLKIEEEEESEAGTVLKQSLPEGTTYDLSKATQ IVLTV	503
consensus	*****	

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NTDB id 20079 SSU05 0428 ABP89395.1	KKATTIQLGNYIGRNSTEVISE	585
NTDB id 146 SP RS08570 WP 000614538.1	KKATTIQLGNYIGRNSTEVISE	526
NTDB id 216 SPD RS08205 WP 000614552.1	KKATTIQLGNYIGRNSTEVISE	526
NTDB id 182 SPR RS07820 WP 000614552.1	KKATTIQLGNYIGRNSTEVISE	526
NTDB id 257 KZH43 RS07655 WP 220041236.1	KKATTIQLGNYIGRNSTEVISE	526
NTDB id 467 HSISS4 RS06915 WP 021143821.1	KKATTIQLGNYIGRNSTEVISE	527
NTDB id 384 SMU RS02325 WP 002263039.1	KKATTIQLGNYIGRNSTEVISE	522
consensus	*****	

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NTDB id 20079 SSU05 0428 ABP89395.1	KKATTIQLGNYIGRNSTEVISE	659
NTDB id 146 SP RS08570 WP 000614538.1	KKATTIQLGNYIGRNSTEVISE	597
NTDB id 216 SPD RS08205 WP 000614552.1	KKATTIQLGNYIGRNSTEVISE	597
NTDB id 182 SPR RS07820 WP 000614552.1	KKATTIQLGNYIGRNSTEVISE	597
NTDB id 257 KZH43 RS07655 WP 220041236.1	KKATTIQLGNYIGRNSTEVISE	597
NTDB id 467 HSISS4 RS06915 WP 021143821.1	KKATTIQLGNYIGRNSTEVISE	598
NTDB id 384 SMU RS02325 WP 002263039.1	KKATTIQLGNYIGRNSTEVISE	596
consensus	*****	

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☒ non conserved
☒ similar
☒ ≥ 50% conserved