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	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	
NTDB id 372 SMU RS05280 WP 002262198.1	0
NTDB id 1419 SGL RS12550 WP 010873252.1	MTSDPNPAQAFENWFEEEEGDDAFTSLIAGEETNSGIEELDFGDLFEQEPTVIQEGEGFGSPSSGEEDPPVRMTLDFNSPE	80
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 35866 SYN6312 RS00135 WP 015122829.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

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	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEIQDLLGVGDVEPAITNPGL	
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NTDB id 1419 SGL RS12550 WP 010873252.1	AELDALFTDEDAPTTAIDPFESSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEIQDLLGVGDVEPAITNPGL	160
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 35866 SYN6312 RS00135 WP 015122829.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

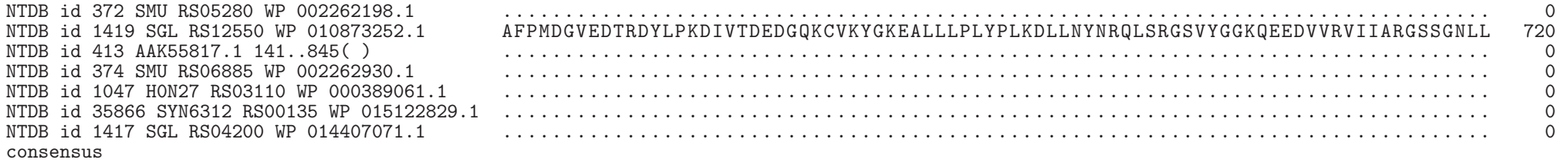
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	HSFAGADRVTPLTNEVLLFAAIDDLGLINAPVLDEEFSFTELETIDTPAPKSFATISHPFRLPVAMEEDEDEFKDL	
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NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
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NTDB id 1417 SGL RS04200 WP 014407071.1	0
consensus		

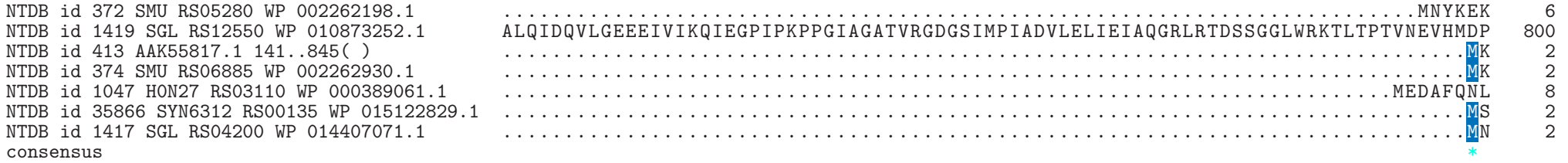
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	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLNLI GELVVKRNRLEQDQERLRQFLDNLNQAQ	
NTDB id 372 SMU RS05280 WP 002262198.1	0
NTDB id 1419 SGL RS12550 WP 010873252.1	LLEQADSSLGGGIAGMGGARLPARPRQVKSRSFEQTMRVPIKQLDNLNLI GELVVKRNRLEQDQERLRQFLDNLNQAQ	320
NTDB id 413 AAK55817.1 141..845()	0
NTDB id 374 SMU RS06885 WP 002262930.1	0
NTDB id 1047 HON27 RS03110 WP 000389061.1	0
NTDB id 35866 SYN6312 RS00135 WP 015122829.1	0
NTDB id 1417 SGL RS04200 WP 014407071.1	0
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 consensus



⊠ non conserved
 ⊠ similar
 ⊠ ≥ 50% conserved