

logo

NTDB id 372 SMU RS05280 WP 002262198.1 ..... 0  
 NTDB id 1419 SGL RS12550 WP 010873252.1 ..... 0  
 NTDB id 413 AAK55817.1 141..845( ) ..... 0  
 NTDB id 374 SMU RS06885 WP 002262930.1 ..... 0  
 NTDB id 1046 HON27 RS03130 WP 168726941.1 MKEILKTLTEAMTLPEDSYSEQDAEFLEIFIEEIEEIEFVDLQPLINKWMQSENIATLTEIRRHFHFTLKGSGRMIGAKSSA 80  
 NTDB id 1047 HON27 RS03110 WP 000389061.1 ..... 0  
 NTDB id 110020 ABXS88 RS01590 WP 353673454.1 ..... 0  
 NTDB id 1417 SGL RS04200 WP 014407071.1 ..... 0  
 consensus ..... 0

logo

NTDB id 372 SMU RS05280 WP 002262198.1 ..... 0  
 NTDB id 1419 SGL RS12550 WP 010873252.1 ..... 0  
 NTDB id 413 AAK55817.1 141..845( ) ..... 0  
 NTDB id 374 SMU RS06885 WP 002262930.1 ..... 0  
 NTDB id 1046 HON27 RS03130 WP 168726941.1 ELAWTVEDTLNRVINQSLQLTPTIQSYVQLVFKFYFLKLVDFNFKQKRAHTLDFRPLILLGQQLQQQSSLEPALEELLQLS 160  
 NTDB id 1047 HON27 RS03110 WP 000389061.1 ..... 0  
 NTDB id 110020 ABXS88 RS01590 WP 353673454.1 ..... 0  
 NTDB id 1417 SGL RS04200 WP 014407071.1 ..... 0  
 consensus ..... 0

logo

NTDB id 372 SMU RS05280 WP 002262198.1 ..... 0  
 NTDB id 1419 SGL RS12550 WP 010873252.1 ..... 0  
 NTDB id 413 AAK55817.1 141..845( ) ..... 0  
 NTDB id 374 SMU RS06885 WP 002262930.1 ..... 0  
 NTDB id 1046 HON27 RS03130 WP 168726941.1 HTLMAETVTGLELDYIEQDSLAEPIIQATHTEIEINLDETTLTFMEEAEEHLATIHQFLDQELHQYDSYNALIRALHTLR 240  
 NTDB id 1047 HON27 RS03110 WP 000389061.1 ..... 0  
 NTDB id 110020 ABXS88 RS01590 WP 353673454.1 ..... 0  
 NTDB id 1417 SGL RS04200 WP 014407071.1 ..... 0  
 consensus ..... 0

logo

NTDB id 372 SMU RS05280 WP 002262198.1 ..... 0  
 NTDB id 1419 SGL RS12550 WP 010873252.1 ..... 0  
 NTDB id 413 AAK55817.1 141..845( ) ..... 0  
 NTDB id 374 SMU RS06885 WP 002262930.1 ..... 0  
 NTDB id 1046 HON27 RS03130 WP 168726941.1 GSSAMAQVETIFEASTKVEHLFKILLQEELSSHSEEILLQEQYREFVRDSELELLSRYSSSEQLLEARLLQFNQSWDAYVEQ 320  
 NTDB id 1047 HON27 RS03110 WP 000389061.1 ..... 0  
 NTDB id 110020 ABXS88 RS01590 WP 353673454.1 ..... 0  
 NTDB id 1417 SGL RS04200 WP 014407071.1 ..... 0  
 consensus ..... 0

logo

	HGDRTDPLMP SHGLV S QLLQLD V S ELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	0
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	HGDRTDPLMP SHGLV S QLLQLD V S ELLDAELDFEKGIRNEFPDYLERLSEQADLLLQHTHSQAMLGLHEYTSQLKESYEV	400
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

logo

	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVVKQHQSI L EELKLYTQYTPDISNDLPQQDSTSFEP I YN I E P	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	.....	3
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	LLDKPALLQSDYIFEIYQKAHQQLIQLFDALAAGQRVGVVVKQHQSI L EELKLYTQYTPDISNDLPQQDSTSFEP I YN I E P	480
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

logo

	EPN EWATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGFARAFENWYEEENBRALFNSLY	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	DPN.....	28
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	EPEVVATVENFSDSADWAVLGQSVQQDRQYISSTQVNRNFDADLLDIFLEEAEEELLEGLDITDLNIWVGEQENFAALNNLM	560
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

logo

	RYL EHTLKGGANMVQATYLG L I AHELES IYERLIQKQLVVTSDLIDFIRLVQDDLADRLQIMREQQLDYAAPYTI NALKRA	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	AGE.....	31
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	RYLHTLKGGANMVQATYLG L I AHELES IYERLIQKQLVVTSDLIDFIRLVQDDLADRLQIMREQQLDYAAPYTI NALKRA	640
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

logo

	<b>E</b> NSN <b>F</b> EE <b>L</b> BY <b>Y</b> BA <b>F</b> BA <b>E</b> PE <b>V</b> F <b>S</b> EE <b>K</b> Y <b>L</b> SE <b>I</b> TT <b>I</b> DE <b>I</b> P <b>V</b> E <b>L</b> E <b>P</b> ALAE <b>L</b> E <b>T</b> H <b>H</b> D <b>Q</b> V <b>F</b> D <b>T</b> AS <b>Z</b> E <b>L</b> ER <b>F</b> P <b>V</b> E <b>M</b> IT <b>S</b> Y <b>K</b> S <b>P</b> E <b>X</b> E <b>Y</b> R	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	ETNSGIEELDFGDLFEGETPTVIQGEFGS.....PSSGEEDPPVRMTLDFNSPEAELD	84
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	GQNSNFQPLPVVEAFDAESEVFSEQKVISEIIDEIPVELEPALAELETHHDQVFDTAVTELATPVEMITTVTSQENEVA	720
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus	.....	0

logo

	<b>A</b> NEQ <b>D</b> IEAW <b>V</b> E <b>L</b> FL <b>B</b> ER <b>T</b> ET <b>T</b> RY <b>M</b> RE <b>L</b> SS <b>S</b> W <b>L</b> E <b>F</b> PS <b>I</b> NP <b>S</b> AD <b>P</b> LA <b>S</b> IP <b>E</b> T <b>N</b> AP <b>A</b> Y <b>Q</b> T <b>A</b> GG <b>L</b> DD <b>N</b> DL <b>A</b> EL <b>Q</b> DL <b>L</b> GV <b>G</b> D <b>V</b> EP	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	.....ALFTDEDAPTTAIDPFSSSSWLEIDQMMEADPLASIPETNAPAYQTAGGLDDNDLAEQLDQLLGVGDVEP	153
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	ANEQDIEAVVEQTFLEEATELLEMAESLLKQWFEQRTNR.....S.....ILL.....	763
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus	.....	0

logo

	<b>A</b> TR <b>B</b> AY <b>H</b> SH <b>K</b> GA <b>R</b> MY <b>E</b> LE <b>N</b> Y <b>G</b> AT <b>A</b> Y <b>A</b> LE <b>N</b> A <b>F</b> E <b>Q</b> F <b>A</b> IB <b>B</b> F <b>B</b> SE <b>N</b> I <b>K</b> RY <b>L</b> BE <b>A</b> F <b>A</b> Y <b>L</b> K <b>D</b> A <b>I</b> F <b>N</b> H <b>N</b> Y <b>Q</b> H <b>F</b> EL <b>L</b> Y <b>S</b> EP <b>A</b> R <b>K</b> S	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	AITNPLGHSFAGADRVTPLTNEVLL.FA.....AIDDLGLINAPVLDEEFSF.....TELETLIDTPAPKS	214
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	.QLQRAVHSLKGGARMVGLEAVQAIAYQLENAFEQFALHHFNSNIYDHLLESIAIWLKDAIFNHNYQHFDGLQQSLENIQ	842
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus	.....	0

logo

	<b>F</b> AT <b>L</b> ET <b>T</b> LS <b>I</b> PY <b>K</b> ME <b>R</b> E <b>D</b> EL <b>F</b> RL <b>E</b> R <b>A</b> LL <b>E</b> Q <b>A</b> D <b>S</b> S <b>L</b> GG <b>I</b> AG <b>M</b> GG <b>A</b> RL <b>P</b> AR <b>P</b> ... <b>R</b> Q <b>V</b> K <b>S</b> R <b>S</b> F <b>E</b> Q <b>T</b> M <b>R</b> V <b>P</b> I <b>K</b> Q <b>L</b> D <b>N</b> L <b>S</b> N <b>L</b> I	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	FATISHPFRLPVAMEEDEDKDLLEALLEQADSSLGGGIAGMGGARLPARP...RQVKRSFEQTMRVPIKQLDNLSNLI	291
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	F..FETTIQIPTKLTRA.DLFSS.EP.....VMTFIQGDGTEPPPMGAWEQTERLDQNNEMIRVSADLIEKMIDLS	910
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus	.....	0

logo

	<b>CENMYRNRRI+ESDDEBFRFLYEMLENRAQRLSDSLFRVMEELYERSTLECALLASRSTNSRHHSSKTTYPDARIAKRSIEH</b>	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	GELVVKRNRLREQDQERLRQFLDNLLNQAQNLSDVGGRMQDLYERTLLEGALLASRSTNSRHHSSKTTYPDARITASSSST	371
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	GENSINRSRIEMDLSQFSHTLVEMELAIQRLADQLRRMEGELETQI.....IAKHGIEH	964
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

logo

	<b>SRYSBDALEMDSYSTHNLSSREMAEAYBYKSPKSTSYKYBETESYABSPSYCTEYQEGMTRIRVVPFISTLABRLPR</b>	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	MNDQPLDALEMDRFTGFHLLSQEMIELIVRVRESASDIQYLVDETDQVARTLRQVTTQLQEGMTKSRMVPFTQTADRLPR	451
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	SRYTDFDPLEMDQYSSLNQLSKSLAESASDLVDFKNTLSDKIRDTESELLLQQSRIQAEIQEGLMRTRLVPPSRLLPRLQR	1044
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

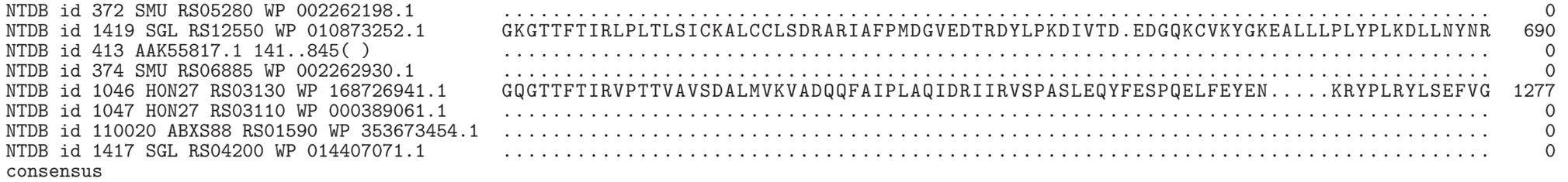
logo

	<b>AYREYSJKLNREXPLEVNNBSEY+DRMILERLYRPMETHLYRNAI+HGH+ESBAERQRAN+KPEEGH+AYRAPEQGNBYMYNY</b>	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	AIREVSMKLNKKVNLEVEGGDVLIDKMLEHLYNPMTHLVNNAITHGIETPAERQKLGKSPGHLAVRAFLQGNQTLITV	531
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	IVRQTSTALNRPAELFVNNTGELDRNILERLVTPLEHMLRNAIDHGLEDRAQRQQANKPETGRIELNIQRQGTDVVVVF	1124
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

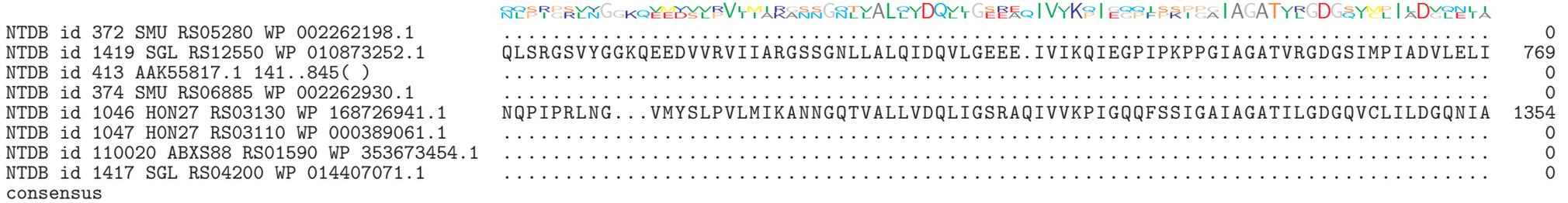
logo

	<b>SDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVVRTSLIDIRGAINIESTV</b>	
NTDB id 372 SMU RS05280 WP 002262198.1	.....	0
NTDB id 1419 SGL RS12550 WP 010873252.1	SDDGAGINTEMVKQKAVEKGLISRMEAQSLSQSEVYDLLFHAGFSTKDTADDFAGRGVGLDVVVRTSLIDIRGAINIESTV	611
NTDB id 413 AAK55817.1 141..845( )	.....	0
NTDB id 374 SMU RS06885 WP 002262930.1	.....	0
NTDB id 1046 HON27 RS03130 WP 168726941.1	SDDGQGIDVEKVRQKALLAGLIKP..EQDLEQQDILQLIFHPGLSTAEQVTQISGRGVGLDVVQSDIKSLGGHVSVESVY	1202
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	0
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	0
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	0
consensus		

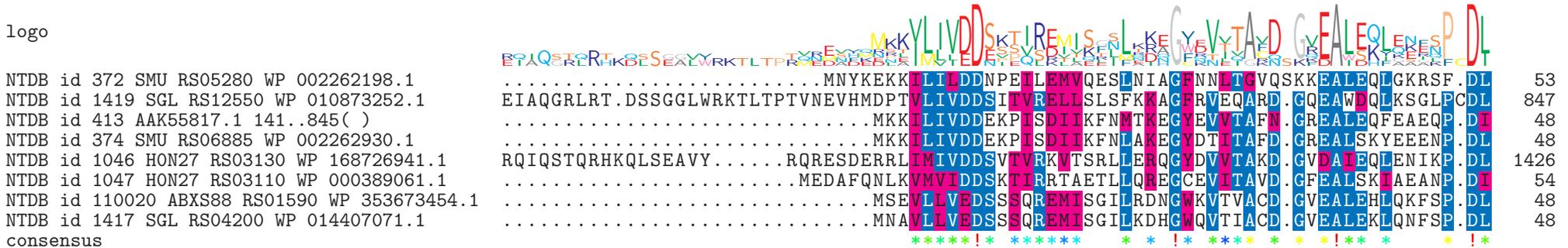
logo



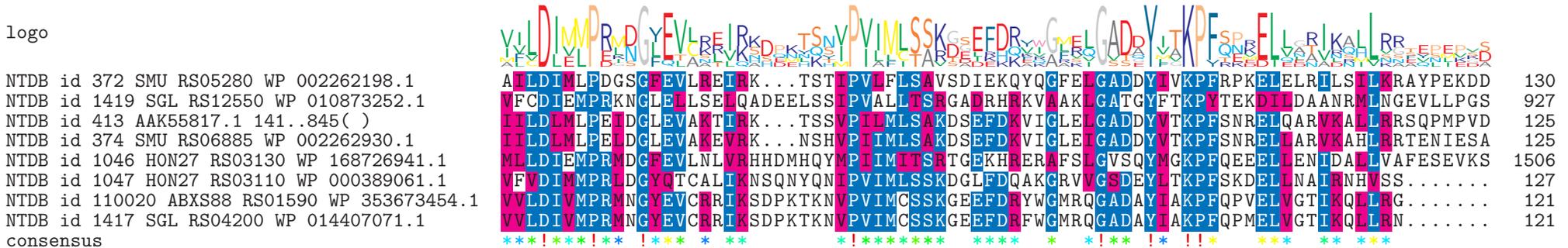
logo



logo



logo



logo

NTDB id 372 SMU RS05280 WP 002262198.1	TLILPTCQVN.....FSQ....ALITKGELEIQLTAKEYSILKVLVYNNQNRIVTFDQLLEK.....V	183
NTDB id 1419 SGL RS12550 WP 010873252.1	TRKTNLSILSRFKKARGNFSEEAPTIVRSNPGTL.....R.....STVLIIDDSVMVREMLSMTFKKADYHTETARDGQDA	997
NTDB id 413 AAK55817.1 141..845( )	GQ.EADSKPQ.PIQIGDLEIVPDAYVAKKYGEELDLTHREFELLYHLASHTGQVITREHLLLET.....V	187
NTDB id 374 SMU RS06885 WP 002262930.1	VAEENASGIP.EIIIGDLQILPDAFVAKKRGTEVELTHREFELLHHLATHTGQVMTREHLLLET.....V	188
NTDB id 1046 HON27 RS03130 WP 168726941.1	.....	1506
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	127
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	121
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	121
consensus		

<0

logo

NTDB id 372 SMU RS05280 WP 002262198.1	WGLQYQGYENTMMA.....HIRKIRQKIEANPSK...PECLLTVK.....GLGYKLVKVS.....	229
NTDB id 1419 SGL RS12550 WP 010873252.1	WEKLLKSGLPDCLLILCDIEMPRMNGLELLERIQEDPNLSRIPVAMITSRGAQKMQRMAAEKGAQGYFVKPYIEETLLEASQ	1077
NTDB id 413 AAK55817.1 141..845( )	WGYDYFG...DVRTVDVTV.....RRLREKIEDTPSR...PEYILT.....RRGVGYMRNNA*.....	234
NTDB id 374 SMU RS06885 WP 002262930.1	WGYDYFG...DVRTVDVTV.....RRLREKIEDTPSR...PEYILT.....RRGVGYMKSVD.....	235
NTDB id 1046 HON27 RS03130 WP 168726941.1	.....	1506
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	127
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	121
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	121
consensus		

logo

NTDB id 372 SMU RS05280 WP 002262198.1	.....	229
NTDB id 1419 SGL RS12550 WP 010873252.1	RLIRGEVLLDLAALSAD E	1095
NTDB id 413 AAK55817.1 141..845( )	.....	234
NTDB id 374 SMU RS06885 WP 002262930.1	.....	235
NTDB id 1046 HON27 RS03130 WP 168726941.1	.....	1506
NTDB id 1047 HON27 RS03110 WP 000389061.1	.....	127
NTDB id 110020 ABXS88 RS01590 WP 353673454.1	.....	121
NTDB id 1417 SGL RS04200 WP 014407071.1	.....	121
consensus		

- non conserved
- similar
- ≥ 50% conserved