

logo

		MSXLRITCDRLGAAALLDQGLDRLQRALERHREVGGLAEVLYDSDQYERRIASXIEERFGIPLVLRVSPKXK	
NTDB id 1028	TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGLAEVLYDMDGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306	DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLDIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
NTDB id 1227	CJJ81176 RS07050 WP 011812800.1	.....	0
NTDB id 1410	SGL RS13950 WP 010873519.1	.....	0
NTDB id 1355	HI 0298 AAC21962.1	.....	0
NTDB id 1366	NTHI RS01950 WP 011271972.1	.....	0
NTDB id 1384	A4U84 RS08525 WP 010787006.1	.....	0
NTDB id 1468	ASU2 RS05090 WP 014991734.1	.....	0
NTDB id 97404	AGG09 RS06685 WP 004145935.1	.....	0
NTDB id 1436	BW25113 RS00530 WP 001025145.1	.....	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1293	VP RS12240 WP 005479695.1	.....	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	.....	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	.....	0
consensus			

logo

		REYKAKELKAIKALPFPALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	
NTDB id 1028	TT RS08215 WP 011173991.1	ALLPAEKAKELKAIKALPFPALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLEIVVPSGA	158
NTDB id 1306	DR RS10060 WP 027479822.1	RSIRPQTALNLQAFPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGA	157
NTDB id 1227	CJJ81176 RS07050 WP 011812800.1	.....	0
NTDB id 1410	SGL RS13950 WP 010873519.1	.....	0
NTDB id 1355	HI 0298 AAC21962.1	.....	0
NTDB id 1366	NTHI RS01950 WP 011271972.1	.....	0
NTDB id 1384	A4U84 RS08525 WP 010787006.1	.....	0
NTDB id 1468	ASU2 RS05090 WP 014991734.1	.....	0
NTDB id 97404	AGG09 RS06685 WP 004145935.1	.....	0
NTDB id 1436	BW25113 RS00530 WP 001025145.1	.....	0
NTDB id 1168	A1552VC RS11070 WP 000957200.1	.....	0
NTDB id 1401	DSB67 RS12665 WP 010643259.1	.....	0
NTDB id 1293	VP RS12240 WP 005479695.1	.....	0
NTDB id 1110	NGFG RS09235 WP 003689817.1	.....	0
NTDB id 1251	GCO85 RS07725 WP 011946523.1	.....	0
NTDB id 1018	ACIAD RS01685 WP 004920473.1	.....	0
NTDB id 1058	ABD1 RS01615 WP 001274986.1	.....	0
consensus			

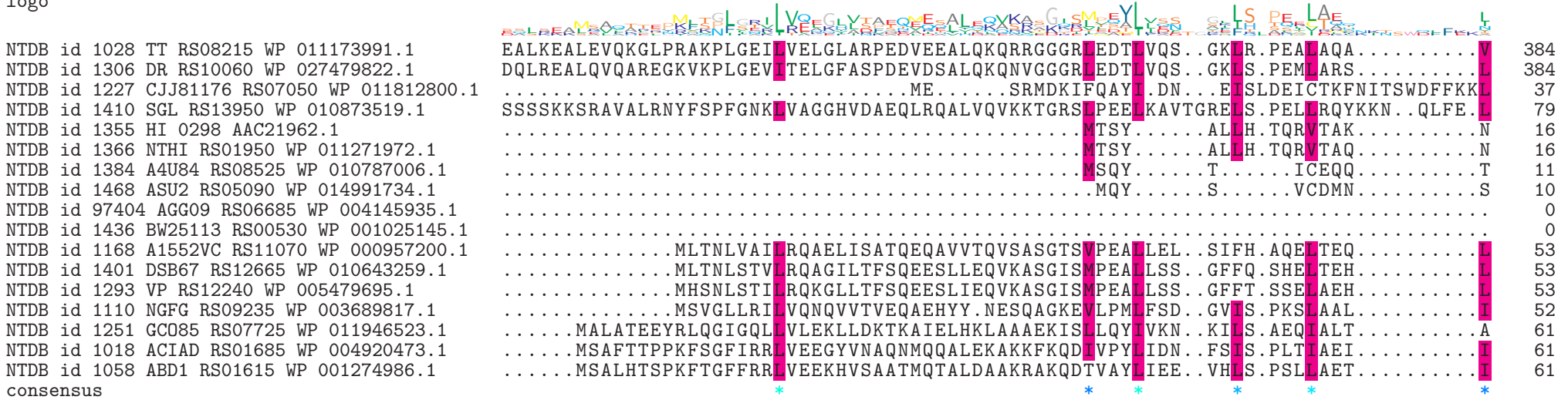
logo

				SRGSGKSLKLEELVGHGXIISKLLYALSKQRTGSKLGRITLYSRKAVTEKLYEXLAEQSGTFLKSTEGAMPSEEXK		
NTDB id 1028	TT	RS08215	WP	011173991.1	SGEGQKDLKLGELLQKQGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSAA	237
NTDB id 1306	DR	RS10060	WP	027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237
NTDB id 1227	CJJ81176	RS07050	WP	011812800.1	.....	0
NTDB id 1410	SGL	RS13950	WP	010873519.1	.....	0
NTDB id 1355	HI	0298	AAC21962.1	.....	0	
NTDB id 1366	NTHI	RS01950	WP	011271972.1	.....	0
NTDB id 1384	A4U84	RS08525	WP	010787006.1	.....	0
NTDB id 1468	ASU2	RS05090	WP	014991734.1	.....	0
NTDB id 97404	AGG09	RS06685	WP	004145935.1	.....	0
NTDB id 1436	BW25113	RS00530	WP	001025145.1	.....	0
NTDB id 1168	A1552VC	RS11070	WP	000957200.1	.....	0
NTDB id 1401	DSB67	RS12665	WP	010643259.1	.....	0
NTDB id 1293	VP	RS12240	WP	005479695.1	.....	0
NTDB id 1110	NGFG	RS09235	WP	003689817.1	.....	0
NTDB id 1251	GCO85	RS07725	WP	011946523.1	.....	0
NTDB id 1018	ACIAD	RS01685	WP	004920473.1	.....	0
NTDB id 1058	ABD1	RS01615	WP	001274986.1	.....	0
consensus						0

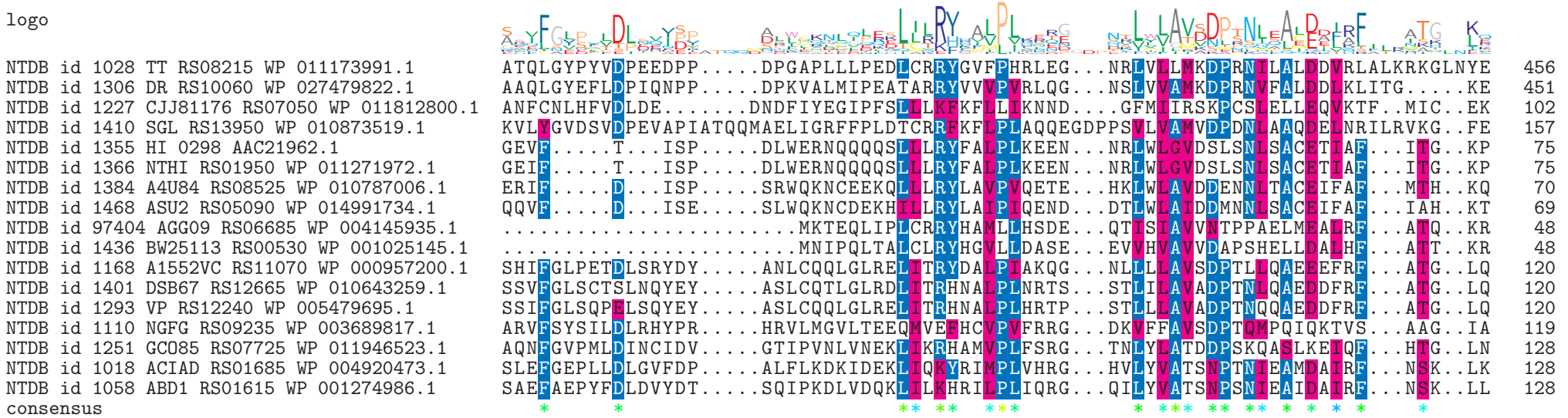
logo

					LEELRSDALRYAVPYSEVVEVYSDPRHKEEVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGQLSR	
NTDB id 1028	TT	RS08215	WP	011173991.1	LLLLRSDALRYGAVPIGFQNGEVEVVLSDPRHKEEVAQLLNRPARFYALALPQAWHEELFRRAYPQKNRLGEVLVQEGQLSR	317
NTDB id 1306	DR	RS10060	WP	027479822.1	GSMLRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQRGRGLGEQLVQEGTSLR	317
NTDB id 1227	CJJ81176	RS07050	WP	011812800.1	.....	0
NTDB id 1410	SGL	RS13950	WP	010873519.1	.....	0
NTDB id 1355	HI	0298	AAC21962.1	.....	MTS	3
NTDB id 1366	NTHI	RS01950	WP	011271972.1	.....	0
NTDB id 1384	A4U84	RS08525	WP	010787006.1	.....	0
NTDB id 1468	ASU2	RS05090	WP	014991734.1	.....	0
NTDB id 97404	AGG09	RS06685	WP	004145935.1	.....	0
NTDB id 1436	BW25113	RS00530	WP	001025145.1	.....	0
NTDB id 1168	A1552VC	RS11070	WP	000957200.1	.....	0
NTDB id 1401	DSB67	RS12665	WP	010643259.1	.....	0
NTDB id 1293	VP	RS12240	WP	005479695.1	.....	0
NTDB id 1110	NGFG	RS09235	WP	003689817.1	.....	0
NTDB id 1251	GCO85	RS07725	WP	011946523.1	.....	0
NTDB id 1018	ACIAD	RS01685	WP	004920473.1	.....	0
NTDB id 1058	ABD1	RS01615	WP	001274986.1	.....	0
consensus						0

logo

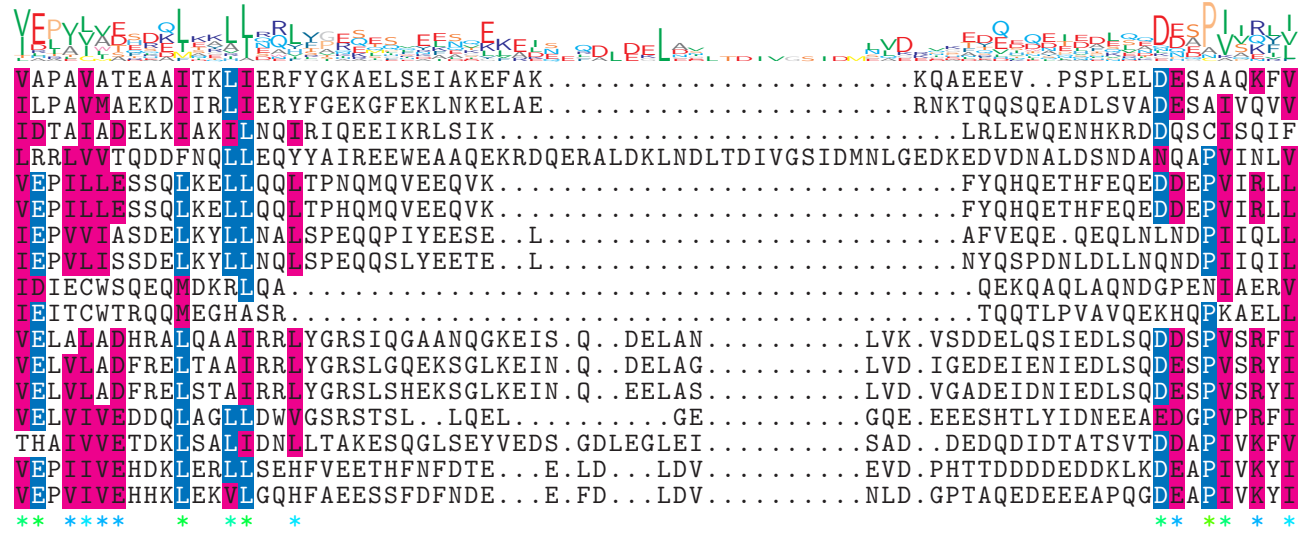


logo



logo

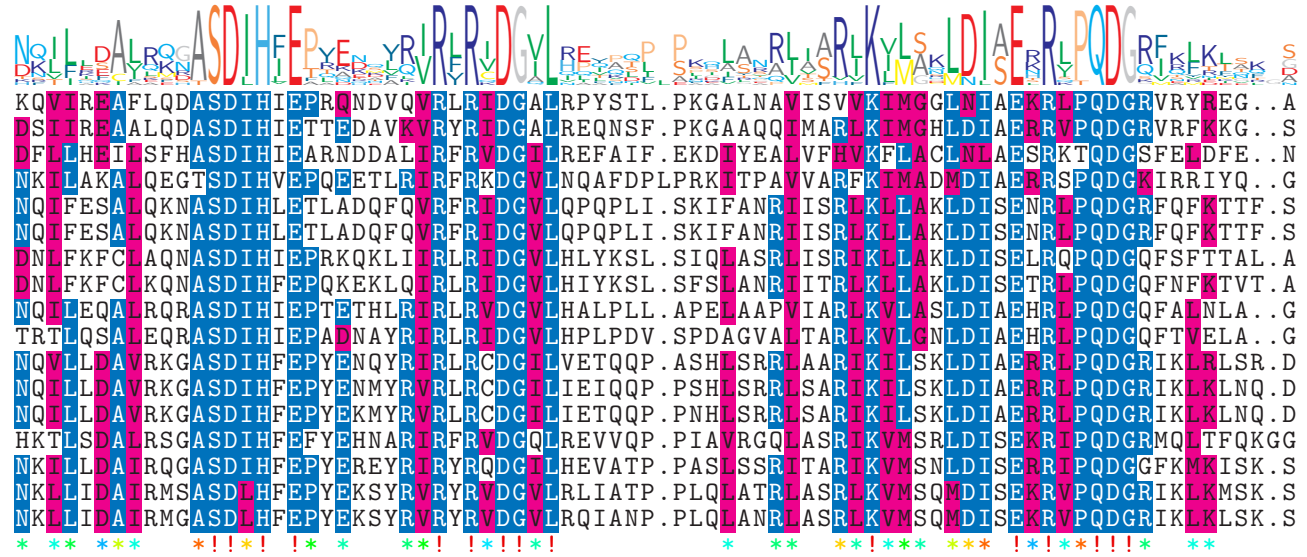
NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
 NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1355 HI 0298 AAC21962.1  
 NTDB id 1366 NTHI RS01950 WP 011271972.1  
 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
 NTDB id 1468 ASU2 RS05090 WP 014991734.1  
 NTDB id 97404 AGG09 RS06685 WP 004145935.1  
 NTDB id 1436 BW25113 RS00530 WP 001025145.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
 NTDB id 1401 DSB67 RS12665 WP 010643259.1  
 NTDB id 1293 VP RS12240 WP 005479695.1  
 NTDB id 1110 NGFG RS09235 WP 003689817.1  
 NTDB id 1251 GC085 RS07725 WP 011946523.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus



511  
508  
153  
237  
126  
126  
121  
121  
85  
85  
186  
186  
186  
176  
195  
190  
190

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
 NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1355 HI 0298 AAC21962.1  
 NTDB id 1366 NTHI RS01950 WP 011271972.1  
 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
 NTDB id 1468 ASU2 RS05090 WP 014991734.1  
 NTDB id 97404 AGG09 RS06685 WP 004145935.1  
 NTDB id 1436 BW25113 RS00530 WP 001025145.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
 NTDB id 1401 DSB67 RS12665 WP 010643259.1  
 NTDB id 1293 VP RS12240 WP 005479695.1  
 NTDB id 1110 NGFG RS09235 WP 003689817.1  
 NTDB id 1251 GC085 RS07725 WP 011946523.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1  
 consensus





logo

NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
 NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1355 HI 0298 AAC21962.1  
 NTDB id 1366 NTHI RS01950 WP 011271972.1  
 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
 NTDB id 1468 ASU2 RS05090 WP 014991734.1  
 NTDB id 97404 AGG09 RS06685 WP 004145935.1  
 NTDB id 1436 BW25113 RS00530 WP 001025145.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
 NTDB id 1401 DSB67 RS12665 WP 010643259.1  
 NTDB id 1293 VP RS12240 WP 005479695.1  
 NTDB id 1110 NGFG RS09235 WP 003689817.1  
 NTDB id 1251 GC085 RS07725 WP 011946523.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1



QATTRLIDEMGV<sup>1</sup>EPFN<sup>1</sup>ISAAL<sup>1</sup>IGVLS<sup>1</sup>QRLV<sup>1</sup>RRV<sup>1</sup>CEH<sup>1</sup>CK<sup>1</sup>VEV.KPDPETLRRLL<sup>1</sup>GLSE.....AE<sup>1</sup>IQGA..RLYK.. 812  
 GAVTRLEEMG<sup>1</sup>IENFN<sup>1</sup>ISAAM<sup>1</sup>MGVLA<sup>1</sup>AQLV<sup>1</sup>RRV<sup>1</sup>CSE<sup>1</sup>CK<sup>1</sup>QPT.NADPEVLRRL<sup>1</sup>GLISE.....RD<sup>1</sup>IRGA..NLMR.. 809  
 STIDRL<sup>1</sup>LDMQAKSY<sup>1</sup>LIASALS<sup>1</sup>LVI<sup>1</sup>AQLV<sup>1</sup>RKLC<sup>1</sup>PWC<sup>1</sup>CK<sup>1</sup>QSKKKHYIE.....FEG.....EFFE... 443  
 GAIARLD<sup>1</sup>EMGV<sup>1</sup>EPFM<sup>1</sup>VSGS<sup>1</sup>LLG<sup>1</sup>VLA<sup>1</sup>QRL<sup>1</sup>MRK<sup>1</sup>VCTE<sup>1</sup>CH<sup>1</sup>IAY.HPSKEELSRF<sup>1</sup>GLSASGDEVVTFYKANS<sup>1</sup>SPDEIHLARQK 554  
 SAISRLL<sup>1</sup>QLG<sup>1</sup>IQQ<sup>1</sup>YE<sup>1</sup>IKNS<sup>1</sup>LLLVIA<sup>1</sup>AQLV<sup>1</sup>RKLC<sup>1</sup>SK<sup>1</sup>CGGN..... 403  
 SAISRLL<sup>1</sup>QLG<sup>1</sup>IQQ<sup>1</sup>HE<sup>1</sup>IENS<sup>1</sup>LLLVIA<sup>1</sup>AQLV<sup>1</sup>RK<sup>1</sup>CPK<sup>1</sup>CGGN..... 403  
 SAVERLL<sup>1</sup>QLG<sup>1</sup>IQE<sup>1</sup>YE<sup>1</sup>IQNS<sup>1</sup>LLLVIA<sup>1</sup>AQLV<sup>1</sup>RKLC<sup>1</sup>KK<sup>1</sup>CGV..... 398  
 SAIERLL<sup>1</sup>QLG<sup>1</sup>IKEYE<sup>1</sup>IKNS<sup>1</sup>LLLVIA<sup>1</sup>AQLV<sup>1</sup>RK<sup>1</sup>CP<sup>1</sup>CGGK..... 398  
 ETILTRL<sup>1</sup>QCMG<sup>1</sup>IARW<sup>1</sup>MISS<sup>1</sup>ALLVIA<sup>1</sup>AQLV<sup>1</sup>RKLC<sup>1</sup>PH<sup>1</sup>CH<sup>1</sup>RNAGSAAD.....LP.....HS<sup>1</sup>WPRPLPRWQ.. 381  
 ETLVRL<sup>1</sup>QCMG<sup>1</sup>VARW<sup>1</sup>MISS<sup>1</sup>ALLVIA<sup>1</sup>AQLV<sup>1</sup>RKLC<sup>1</sup>PH<sup>1</sup>CH<sup>1</sup>RQQGGEPIH.....IP.....DN<sup>1</sup>WPSPLPHWQ.. 381  
 ETVIRLAN<sup>1</sup>MGVE<sup>1</sup>PFNL<sup>1</sup>ASS<sup>1</sup>LSL<sup>1</sup>IA<sup>1</sup>AQL<sup>1</sup>ARR<sup>1</sup>LCK<sup>1</sup>HCK<sup>1</sup>IAV.RPSALLQSQF<sup>1</sup>AFQP.....NE<sup>1</sup>IL....YEAN.. 486  
 ETVIRLS<sup>1</sup>NMGV<sup>1</sup>ESFNL<sup>1</sup>ASS<sup>1</sup>LSL<sup>1</sup>IA<sup>1</sup>AQL<sup>1</sup>ARK<sup>1</sup>LC<sup>1</sup>PY<sup>1</sup>CH<sup>1</sup>QPPQ.EPNSQLQH.IGIAN.....TEQI....FQAN.. 485  
 ETVIRLS<sup>1</sup>NMGV<sup>1</sup>ESFNL<sup>1</sup>ASS<sup>1</sup>LSL<sup>1</sup>IA<sup>1</sup>AQL<sup>1</sup>ARK<sup>1</sup>LC<sup>1</sup>PY<sup>1</sup>CK<sup>1</sup>QPPQ.EHTVQLQH.LGIQT.....TDNI....FRAN.. 485  
 ATLSR<sup>1</sup>MLNMGV<sup>1</sup>APFN<sup>1</sup>IAS<sup>1</sup>SV<sup>1</sup>SL<sup>1</sup>MA<sup>1</sup>AQL<sup>1</sup>LR<sup>1</sup>RL<sup>1</sup>CSS<sup>1</sup>CK<sup>1</sup>QEVERPSASALKEVGF<sup>1</sup>TD.....ED<sup>1</sup>LAK.DWKLYR.. 481  
 ETLNRL<sup>1</sup>VNMG<sup>1</sup>IPTFN<sup>1</sup>IAS<sup>1</sup>SV<sup>1</sup>TL<sup>1</sup>IA<sup>1</sup>AQL<sup>1</sup>ARK<sup>1</sup>LN<sup>1</sup>QCK<sup>1</sup>AVRDDFTN<sup>1</sup>QGLIEL<sup>1</sup>GFKE.....AD<sup>1</sup>V..NLKLYK.. 498  
 ETLTRL<sup>1</sup>RNMGV<sup>1</sup>ASFNI<sup>1</sup>ATS<sup>1</sup>VN<sup>1</sup>LVIA<sup>1</sup>AQL<sup>1</sup>ARR<sup>1</sup>LS<sup>1</sup>QCK<sup>1</sup>RPI.QVPERSLLEM<sup>1</sup>GF<sup>1</sup>TP.....ED<sup>1</sup>LAQPEFQIFE.. 494  
 ETLTRL<sup>1</sup>RNMGV<sup>1</sup>PSFNI<sup>1</sup>ATS<sup>1</sup>VN<sup>1</sup>LVIA<sup>1</sup>AQL<sup>1</sup>ARR<sup>1</sup>LS<sup>1</sup>QCK<sup>1</sup>IPA.DIPKQSLLEM<sup>1</sup>GF<sup>1</sup>TE.....QD<sup>1</sup>LAHPDFRVFQ.. 494

consensus

\*\*\* !\* \*\*\*\* \* \* \*\*\* \*\*\*\*\*!!\*!\*!!\*!\* !!\* \* \*

logo

NTDB id 1028 TT RS08215 WP 011173991.1  
 NTDB id 1306 DR RS10060 WP 027479822.1  
 NTDB id 1227 CJJ81176 RS07050 WP 011812800.1  
 NTDB id 1410 SGL RS13950 WP 010873519.1  
 NTDB id 1355 HI 0298 AAC21962.1  
 NTDB id 1366 NTHI RS01950 WP 011271972.1  
 NTDB id 1384 A4U84 RS08525 WP 010787006.1  
 NTDB id 1468 ASU2 RS05090 WP 014991734.1  
 NTDB id 97404 AGG09 RS06685 WP 004145935.1  
 NTDB id 1436 BW25113 RS00530 WP 001025145.1  
 NTDB id 1168 A1552VC RS11070 WP 000957200.1  
 NTDB id 1401 DSB67 RS12665 WP 010643259.1  
 NTDB id 1293 VP RS12240 WP 005479695.1  
 NTDB id 1110 NGFG RS09235 WP 003689817.1  
 NTDB id 1251 GC085 RS07725 WP 011946523.1  
 NTDB id 1018 ACIAD RS01685 WP 004920473.1  
 NTDB id 1058 ABD1 RS01615 WP 001274986.1



GMGCE..RCGGT<sup>1</sup>G<sup>1</sup>YK<sup>1</sup>G<sup>1</sup>RYA<sup>1</sup>THE<sup>1</sup>LLV<sup>1</sup>VDDE<sup>1</sup>IRHA<sup>1</sup>TVAGKSATE<sup>1</sup>KEI<sup>1</sup>AR.RK<sup>1</sup>CM<sup>1</sup>K<sup>1</sup>TL<sup>1</sup>RED<sup>1</sup>GL<sup>1</sup>YK<sup>1</sup>AL<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>IT<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>LARTIE 889  
 GTGCP..RCGGT<sup>1</sup>G<sup>1</sup>YK<sup>1</sup>GR<sup>1</sup>M<sup>1</sup>G<sup>1</sup>IHE<sup>1</sup>LM<sup>1</sup>VDDSL<sup>1</sup>RRT<sup>1</sup>IGAGRPAAE<sup>1</sup>IRD<sup>1</sup>VALGES<sup>1</sup>CL<sup>1</sup>RS<sup>1</sup>LR<sup>1</sup>QD<sup>1</sup>GL<sup>1</sup>EK<sup>1</sup>AL<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>IT<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>LAV<sup>1</sup>TAS 887  
 PKGCE..RCHHS<sup>1</sup>CF<sup>1</sup>FG<sup>1</sup>REL<sup>1</sup>IAB<sup>1</sup>CL<sup>1</sup>EINED<sup>1</sup>LACAT<sup>1</sup>RENQDKT<sup>1</sup>ILMELAK.KY<sup>1</sup>CF<sup>1</sup>Q<sup>1</sup>TM<sup>1</sup>FEG<sup>1</sup>CL<sup>1</sup>KK<sup>1</sup>AKE<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>IDE<sup>1</sup>LLRV<sup>1</sup>R. 519  
 GNLCQ..KCSGS<sup>1</sup>G<sup>1</sup>YK<sup>1</sup>GR<sup>1</sup>V<sup>1</sup>G<sup>1</sup>YEV<sup>1</sup>MRNSER<sup>1</sup>IAEL<sup>1</sup>INQ<sup>1</sup>GATTDR<sup>1</sup>IKDCAV.EEC<sup>1</sup>M<sup>1</sup>IT<sup>1</sup>LLAYS<sup>1</sup>LN<sup>1</sup>VQ<sup>1</sup>E<sup>1</sup>GY<sup>1</sup>IT<sup>1</sup>LEE<sup>1</sup>VER<sup>1</sup>V<sup>1</sup>TFT 631  
 ..LANSCDCH.QGYRGR<sup>1</sup>IG<sup>1</sup>VY<sup>1</sup>Q<sup>1</sup>FL<sup>1</sup>HWQQN.....DY.....Q<sup>1</sup>TD<sup>1</sup>FK<sup>1</sup>NL<sup>1</sup>RA<sup>1</sup>SG<sup>1</sup>EK<sup>1</sup>V<sup>1</sup>SQ<sup>1</sup>GL<sup>1</sup>T<sup>1</sup>DEKE<sup>1</sup>IERV<sup>1</sup>IGK 462  
 ..LINS<sup>1</sup>CDCH.QGYRGR<sup>1</sup>IG<sup>1</sup>VY<sup>1</sup>Q<sup>1</sup>FL<sup>1</sup>HWQQN.....GY.....Q<sup>1</sup>TD<sup>1</sup>FN<sup>1</sup>LR<sup>1</sup>ES<sup>1</sup>GL<sup>1</sup>EK<sup>1</sup>V<sup>1</sup>SQ<sup>1</sup>GL<sup>1</sup>T<sup>1</sup>DEKE<sup>1</sup>IERV<sup>1</sup>IGK 462  
 ..GCD..ECY.QGYRGR<sup>1</sup>IG<sup>1</sup>VY<sup>1</sup>Q<sup>1</sup>CL<sup>1</sup>SRTAK.....K<sup>1</sup>FD<sup>1</sup>KQ<sup>1</sup>TA.YL<sup>1</sup>DF<sup>1</sup>SS<sup>1</sup>LR<sup>1</sup>ES<sup>1</sup>AK<sup>1</sup>Q<sup>1</sup>KE<sup>1</sup>EQ<sup>1</sup>GL<sup>1</sup>T<sup>1</sup>NETE<sup>1</sup>V<sup>1</sup>DR<sup>1</sup>V<sup>1</sup>IGY 460  
 ..GCD..NCY.HGYKGR<sup>1</sup>T<sup>1</sup>GY<sup>1</sup>QL<sup>1</sup>LAKQ<sup>1</sup>GK.....FFEKSTA.YL<sup>1</sup>DY<sup>1</sup>DN<sup>1</sup>LR<sup>1</sup>AS<sup>1</sup>AE<sup>1</sup>Q<sup>1</sup>KL<sup>1</sup>AAN<sup>1</sup>IT<sup>1</sup>V<sup>1</sup>DE<sup>1</sup>V<sup>1</sup>KR<sup>1</sup>V<sup>1</sup>IGD 460  
 AAGCE..HCY.HGYGR<sup>1</sup>LALF<sup>1</sup>EV<sup>1</sup>LPV<sup>1</sup>TPG<sup>1</sup>IRQ<sup>1</sup>GT<sup>1</sup>VQGL<sup>1</sup>NAIE<sup>1</sup>ES<sup>1</sup>LAR.AAG<sup>1</sup>MM<sup>1</sup>TL<sup>1</sup>FE<sup>1</sup>SG<sup>1</sup>CQ<sup>1</sup>ATE<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>RV<sup>1</sup>LGI 457  
 APGCV..HCY.HGFYGR<sup>1</sup>TALF<sup>1</sup>EV<sup>1</sup>LPIT<sup>1</sup>PV<sup>1</sup>IRQL<sup>1</sup>ISANTD<sup>1</sup>VES<sup>1</sup>LE<sup>1</sup>THAR.QA<sup>1</sup>CM<sup>1</sup>RT<sup>1</sup>LFEN<sup>1</sup>CL<sup>1</sup>AVE<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>FE<sup>1</sup>EL<sup>1</sup>IRV<sup>1</sup>LGM 457  
 AAGCN..ECT.GGYSGR<sup>1</sup>V<sup>1</sup>GYEV<sup>1</sup>MAFN<sup>1</sup>TE<sup>1</sup>LAEAT<sup>1</sup>MQRAS<sup>1</sup>IHQ<sup>1</sup>TERLAK.AN<sup>1</sup>CM<sup>1</sup>QT<sup>1</sup>LQES<sup>1</sup>GL<sup>1</sup>EK<sup>1</sup>LR<sup>1</sup>E<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>FAE<sup>1</sup>LQR<sup>1</sup>V<sup>1</sup>LYF 562  
 PDGCN..ECT.HGYSGR<sup>1</sup>T<sup>1</sup>GYEV<sup>1</sup>MK<sup>1</sup>FDES<sup>1</sup>SEAL<sup>1</sup>IKGASV<sup>1</sup>HELE<sup>1</sup>KLAI.AN<sup>1</sup>CM<sup>1</sup>QT<sup>1</sup>LQMS<sup>1</sup>GL<sup>1</sup>EK<sup>1</sup>LK<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>FR<sup>1</sup>EL<sup>1</sup>QR<sup>1</sup>V<sup>1</sup>LYF 561  
 PDGCN..ECT.HGYSGR<sup>1</sup>T<sup>1</sup>GYEV<sup>1</sup>MR<sup>1</sup>FDES<sup>1</sup>SEAL<sup>1</sup>IKGASV<sup>1</sup>HELE<sup>1</sup>KLAI.AN<sup>1</sup>CM<sup>1</sup>ST<sup>1</sup>LQMS<sup>1</sup>GL<sup>1</sup>EK<sup>1</sup>LK<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>FS<sup>1</sup>EL<sup>1</sup>QR<sup>1</sup>V<sup>1</sup>LYF 561  
 AVGCD..RCRGQ<sup>1</sup>G<sup>1</sup>YK<sup>1</sup>GR<sup>1</sup>AG<sup>1</sup>VYEV<sup>1</sup>MP<sup>1</sup>ISEEM<sup>1</sup>QRV<sup>1</sup>IMNNGTE<sup>1</sup>VGL<sup>1</sup>LDVAY.KE<sup>1</sup>CM<sup>1</sup>V<sup>1</sup>DL<sup>1</sup>RR<sup>1</sup>AG<sup>1</sup>IL<sup>1</sup>K<sup>1</sup>IM<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>TANT<sup>1</sup>ND 558  
 AVGCE..QCT.SGYRGR<sup>1</sup>V<sup>1</sup>GLF<sup>1</sup>EV<sup>1</sup>LPMT<sup>1</sup>KE<sup>1</sup>L<sup>1</sup>GQL<sup>1</sup>IMSGGNS<sup>1</sup>LD<sup>1</sup>LKLAQ.SEC<sup>1</sup>ML<sup>1</sup>TL<sup>1</sup>FQ<sup>1</sup>SG<sup>1</sup>EK<sup>1</sup>V<sup>1</sup>KE<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>NR<sup>1</sup>V<sup>1</sup>TD 574  
 PVGCH..DCR.EGYKGR<sup>1</sup>V<sup>1</sup>GYEV<sup>1</sup>MK<sup>1</sup>IT<sup>1</sup>PE<sup>1</sup>ISKI<sup>1</sup>IMEDGNALE<sup>1</sup>IAATAE.TL<sup>1</sup>GF<sup>1</sup>NN<sup>1</sup>LR<sup>1</sup>SG<sup>1</sup>LK<sup>1</sup>V<sup>1</sup>M<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>NR<sup>1</sup>V<sup>1</sup>TSE 570  
 PVGCL..ECR.EGYKGR<sup>1</sup>V<sup>1</sup>GYEV<sup>1</sup>MKV<sup>1</sup>TPE<sup>1</sup>ISKI<sup>1</sup>IMEDGNALE<sup>1</sup>IAAASE.KL<sup>1</sup>GF<sup>1</sup>NN<sup>1</sup>LR<sup>1</sup>SG<sup>1</sup>LK<sup>1</sup>V<sup>1</sup>M<sup>1</sup>Q<sup>1</sup>GL<sup>1</sup>TS<sup>1</sup>LEE<sup>1</sup>V<sup>1</sup>NR<sup>1</sup>V<sup>1</sup>TSE 570

consensus

\*\* ! !\*\*!!\* \*\*\*\* \* \* \*\*\* \*\*\*\*\*!!\*!\*!!\*!\* !!\* \* \*

logo

NTDB id 1028 TT RS08215 WP 011173991.1	.....	889
NTDB id 1306 DR RS10060 WP 027479822.1	.....	887
NTDB id 1227 CJJ81176 RS07050 WP 011812800.1	.....	519
NTDB id 1410 SGL RS13950 WP 010873519.1	DTGLESEMRAKRKKSALTTCRTCAAELQQEWLDCPYCMTSRFS	672
NTDB id 1355 HI 0298 AAC21962.1	NL.....	464
NTDB id 1366 NTHI RS01950 WP 011271972.1	NS.....	464
NTDB id 1384 A4U84 RS08525 WP 010787006.1	E.....	461
NTDB id 1468 ASU2 RS05090 WP 014991734.1	G.....	461
NTDB id 97404 AGG09 RS06685 WP 004145935.1	PHGD.....	461
NTDB id 1436 BW25113 RS00530 WP 001025145.1	PHGE.....	461
NTDB id 1168 A1552VC RS11070 WP 000957200.1	.....	562
NTDB id 1401 DSB67 RS12665 WP 010643259.1	.....	561
NTDB id 1293 VP RS12240 WP 005479695.1	.....	561
NTDB id 1110 NGFG RS09235 WP 003689817.1	.....	558
NTDB id 1251 GC085 RS07725 WP 011946523.1	.....	574
NTDB id 1018 ACIAD RS01685 WP 004920473.1	.....	570
NTDB id 1058 ABD1 RS01615 WP 001274986.1	.....	570
consensus		

DTGLESEMRAKRKKSALTTCRTCAAELQQEWLDCPYCMTSRFS

- non conserved
- similar
- ≥ 50% conserved