

logo

NTDB id 1306 DR RS10060 WP 027479822.1	MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1071543 ACIX90 RS04985 WP 005601209.1		0
NTDB id 1384 A4U84 RS08525 WP 010787006.1		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 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	SIRPQTALNLQAFPPFALEGDRRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLEIVVPSGASDA	160
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1071543 ACIX90 RS04985 WP 005601209.1		0
NTDB id 1384 A4U84 RS08525 WP 010787006.1		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 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

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NTDB id 1306 DR RS10060 WP 027479822.1	GRTGGKLGGERLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1071543 ACIX90 RS04985 WP 005601209.1		0
NTDB id 1384 A4U84 RS08525 WP 010787006.1		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 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

NTDB id 1306 DR RS10060 WP 027479822.1	LRADALRLLAVPVDETEQGVTVLTS DPRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRGLGEQLVQEGTLSRDQL	320
NTDB id 1355 HI 0298 AAC21962.1		0
NTDB id 1366 NTHI RS01950 WP 011271972.1		0
NTDB id 1071543 ACIX90 RS04985 WP 005601209.1		0
NTDB id 1384 A4U84 RS08525 WP 010787006.1		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 1251 GC085 RS07725 WP 011946523.1		0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		0
consensus		

NTDB id 1306	DR RS10060 WP 027479822.1	REALQVQAREGKVKPLGEVITELGFASPDVDSA	QKQNVGGGRLED	LVQSGKLSPEMLARS	AAQLG	GYEFLDPIQNP	400
NTDB id 1355	HI 0298 AAC21962.1			MTSYAL		LHTQRVTA	KN
NTDB id 1366	NTHI RS01950 WP 011271972.1			MTSYAL		LHTQRVTA	QNG
NTDB id 1071543	ACIX90 RS04985 WP 005601209.1			MQY		SVC	DVNNR
NTDB id 1384	A4U84 RS08525 WP 010787006.1			MSQY		TICE	QQT
NTDB id 1168	A1552VC RS11070 WP 000957200.1		MLTNLVAT	LRQAEL	ISATQEQAV	VTQVSASGTS	VPEAL
NTDB id 1401	DSB67 RS12665 WP 010643259.1		MLTNLSTV	LRQAGIL	TFSQEESL	IEQVKASGIS	MPEAL
NTDB id 1293	VP RS12240 WP 005479695.1		MHSNLSTI	LRQKGL	TFSQEESL	IEQVKASGIS	MPEAL
NTDB id 1251	GC085 RS07725 WP 011946523.1	...	MALATEEYRL	QGIGQL	LVLEKLL	DKTKAIEL	HLKAAAEKIS
NTDB id 1058	ABD1 RS01615 WP 001274986.1	...	MSALHTSP	KFTGFFRR	LVEEKHV	SAATMQTA	DAAKRAKQD
NTDB id 1018	ACIAD RS01685 WP 004920473.1	...	MSAFTTPPK	FSGFIRRL	VEEGYV	NAQNMQA	LEKAKFKQD
consensus				*	*	*	*

NTDB	id	1306	DR	RS10060	WP	027479822.1	DPKVALMIPEATARRYVVVPVRLQGNLSLVAMKDPRNVFALDDKLKITGKEILPAVMAEKDITIRLIERFYGEKGFEKLNK	480
NTDB	id	1355	HI	0298	AAC21962.1		DLWERNQQQQSLILRYFALPLKEENNRLLWLGVDLSNLNSACETIAFITGKPVEPILLESSQLKELLQQLTPNQMQVVEEQV	104
NTDB	id	1366	NTHI	RS01950	WP	011271972.1	DLWERNQQQQSLILRYFALPLKEENNRLLWLGVDLSNLNSACETIAFITGKPVEPILLESSQLKELLQQLTPHQMVSVEEQV	104
NTDB	id	1071543	ACIX90	RS04985	WP	005601209.1	EQWQKNYDERHILLRLYLTVPLREDQQTWLAIIDMNNLSACEIFAFLTHKRTEPVVISSEELKYLLGLMPQEQQSVYEED	98
NTDB	id	1384	A4U84	RS08525	WP	010787006.1	SRWQKNCEEKQILLRYLAVPVQETEHLWLAVDDENNLTACEIFAFMTHTKHQIEPVVIASDELKYLINALSPEQQPIYEES	99
NTDB	id	1168	A1552VC	RS11070	WP	000957200.1	ANLCQQIGLRELITRYDALPIAKQGNIALLAVSDPTLLQAEEEFRAFATGLQVELALADHRALQAAITRRLYGRSIQGAANQ	149
NTDB	id	1401	DSB67	RS12665	WP	010643259.1	ASLCQTLGLRDILITRHNALPLNRTSSSTLILAVADPTNLQAEDDFRAFATGLQVELVLADFRELTAATITRRLYGRSLGQEKSG	149
NTDB	id	1293	VP	RS12240	WP	005479695.1	ASLCQQIGLRELITRHNALPLHRTPTSTLLAVADPTNQQAEDDFRAFATGLQVELVLADFRELSTAATITRRLYGRSLSHEKSG	149
NTDB	id	1251	GC085	RS07725	WP	011946523.1	G TIPVN LVNEKL I KRHAM V P LFSRGTNLYLATD DPSKQASLK E T Q F H T G L N T H A I V V E T D K L S A L I D N L L T A K E S Q G L S E	157
NTDB	id	1058	ABD1	RS01615	WP	001274986.1	SQIPKDLVDQKLI L K H R I L P L I Q R G H I Y V A T S N P S N I E A I D A I R F N S K L L V E P V I V E H H K L E K V L G Q H F A E S S F D F N D	157
NTDB	id	1018	ACIAD	RS01685	WP	004920473.1	ALFLKDKIDEKLI Q KYRI M P L V H R G H V L Y V A T S N P T N I E A M D A I R F N S K L K V E P I I V E H D K L E R L I S E H F V E E T H F N F D T	157
consensus							*** * * * * ! * * * * * * * * * * * * * * * *	

