

NTDB id 1402 DSB67 RS12670 WP 010643257.1	MKTIA PQ L K NY R W K G I NS S G K . K T S G N V L A M T E I E V R E R L D A Q H I K I K L K G S I S F I T . K I S	61
NTDB id 1169 A1552VC RS11075 WP 000648511.1	MKAT Q T L P L K N Y R W K G I NS N G K . K V S Q M L A I S E I E V R D K L K D Q H I Q I K L K L G S V S . L A . R L T	62
NTDB id 1113 AAA85695.1 219..1451()	MAK.....NGGF S L F A K K E K R F I E G R H S A S D K L V N G E V S A F T E E E A R K L A K R G I R P L Q I T R V KT S S K	64
NTDB id 1112 NGFG RS09215 WP 003689811.1	MAK.....NGGF S L F A K K E K R F I E G R H S A S D K L V N G E V S A F T E E E A R K L A K R G I R P L Q I T R V KT S S K	64
NTDB id 1252 GC085 RS07730 WP 011213805.1	MD K N S P P L T F H Y Q G I N K A G Q . K M E G D I Q A S R L A I A K A D L R K Q G I V T N K V V K R K P . F D . R K N	61
NTDB id 394636 FM038 RS01910 WP 142871698.1	MATAAT I K R Q T R N K N A K S Q P K V F T F E W K G V N R D G Q . K T S G E L K G S T A A E I R S Q L K S Q G V N P K S V K Q S A S L F K . L G D	78
NTDB id 1198 PSJM300 03950 AFN76868.1	MA Q K A I K N S V F T E G L D R Q G A. K I K G E L S G V S P A L V K A Q L R K Q G I N P Q K V R K S I S L F S .. A G	60
NTDB id 1016 ACIAD RS01680 WP 004920476.1	MA A K T Q V M P V F A Y E G V D R K G I . K L K G E L P A R N M A L A K V T L R K Q G I T I R T I R E K R K N I F E G L L K	63
NTDB id 1059 ABD1 RS01610 WP 000279215.1	MA V K K A Q M P T F A Y E G V D R K G V . K I K G E L P A K N M A L A K V T L R K Q G V T V N I R E K R K N I L E G L F K	63
consensus	* * * * * * * * * *	

[illegible][illegible]

NTDB id 1402	DSB67 RS12670 WP 010643257.1	TQNWGPFGLSFVSLISGKIMSQRSDSFRLLAARSSLRVPILGPVLSKAAIAKFSRTLATSFSAGIPILTALKTTSTKS	301
NTDB id 1169	A1552VC RS11075 WP 000648511.1	VQAYSLWAFIAIAAAIFGLKALRKNSFQIRLKTSRLGLKFPIICNVLA KASIAKFSRTLATSFAAGIPILASLKTAKTS	302
NTDB id 1113	AAA85695.1 219..1451()	FVSYGVWVLIALGFAYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAA	304
NTDB id 1112	NGFG RS09215 WP 003689811.1	FVSYGVWVLIALGFAYRFLKLKARSIKIQRRMDAILLRMPIFGDIVRKGTIARWGRTTATLFAAGVPLVDVLDSTAGAA	304
NTDB id 1252	GCO85 RS07730 WP 011213805.1	MQAYYWITFGALGGVVYSFFYAKNHSLEFAQTIDRVMLKLVPVIGPILEKAAIARFARTLTITFAAGPLVEALKSVAGAT	301
NTDB id 394636	FM038 RS01910 WP 142871698.1	LQASWYIFAVAITIITILFKRAHLHLSQKFRDQVDAAMLKIPAIGPILHKAAAMARFARTLSTTFAAGVPLIDGLESAAGAS	318
NTDB id 1198	PSJM300 03950 AFN76868.1	LQAWWYVVLAGFVCAAYAFKQAHTRSEKFRNWFDRMLMKVPVVGDI LYKSSVARFARTLSTTFAAGVPLVDALDSVAGAT	300
NTDB id 1016	ACIAD RS01680 WP 004920476.1	MQDYWFILIIIVTCAAIAGFLEARKRSKRFRDQLDLRLVRLPIVGD LVYKAIIARYSRTLATTFAAGVPLIDALASTAGAT	303
NTDB id 1059	ABD1 RS01610 WP 000279215.1	MQEYWFIMIIAICAVIAAFLEAKKRSKKFRDGLDKLALKLPVIFGDL VYKAIIARYSRTLATTFAAGVPLIDALESTAGAT	303
consensus		* * * * * ! * * * * * * * * * * ! * * * * * * * * * * ! * * * * * ! * * * * *	

