

logo

		MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	
NTDB id 1306 DR RS10060 WP 027479822.1		MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLLAVQPDPAALR	80
NTDB id 1168 A1552VC RS11070 WP 000957200.1		.....	0
NTDB id 1293 VP RS12240 WP 005479695.1		.....	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1		.....	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		.....	0
NTDB id 365512 IX0704 RS07780 WP 011408072.1		.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		.....	0
consensus			

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		SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEWVPSGASDA	
NTDB id 1306 DR RS10060 WP 027479822.1		SIRPQTALNLQAFPFALEGDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLIEWVPSGASDA	160
NTDB id 1168 A1552VC RS11070 WP 000957200.1		.....	0
NTDB id 1293 VP RS12240 WP 005479695.1		.....	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1		.....	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		.....	0
NTDB id 365512 IX0704 RS07780 WP 011408072.1		.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		.....	0
consensus			

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		GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	
NTDB id 1306 DR RS10060 WP 027479822.1		GRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVLGSM	240
NTDB id 1168 A1552VC RS11070 WP 000957200.1		.....	0
NTDB id 1293 VP RS12240 WP 005479695.1		.....	0
NTDB id 1401 DSB67 RS12665 WP 010643259.1		.....	0
NTDB id 1110 NGFG RS09235 WP 003689817.1		.....	0
NTDB id 1251 GCO85 RS07725 WP 011946523.1		.....	0
NTDB id 365512 IX0704 RS07780 WP 011408072.1		.....	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1		.....	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1		.....	0
consensus			

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		LRADALRLLAVPVDETEQGVTVLTSDFRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRIGEQLVQEGTISRDL	
NTDB id 1306 DR RS10060 WP 027479822.1		LRADALRLLAVPVDETEQGVTVLTSDFRKRPDIDALIGRPVQLMLTRPRDIERLIEQFYPQGRIGEQLVQEGTISRDL	320
NTDB id 1168 A1552VC RS11070 WP 000957200.1		..... MLINLVALRQAEELISATQ	19
NTDB id 1293 VP RS12240 WP 005479695.1		..... MHSNLSLRLRQKGLTFSQ	19
NTDB id 1401 DSB67 RS12665 WP 010643259.1		..... MLNLSLRLRQKGLTFSQ	19
NTDB id 1110 NGFG RS09235 WP 003689817.1		..... MSVGLRLRLVQNQVITVEQ	19
NTDB id 1251 GCO85 RS07725 WP 011946523.1		..... MALATEEYRLQGTGQLVLEKLDKTK	27
NTDB id 365512 IX0704 RS07780 WP 011408072.1		..... MNSVSTTNLVGITARRLVQDGALEETA	29
NTDB id 1058 ABD1 RS01615 WP 001274986.1		..... MS..ALHTSPKFTGFFRRLVEEKHVSAAAT	27
NTDB id 1018 ACIAD RS01685 WP 004920473.1		..... MS..AFTTPPKFSGFIRRLVEEGYVNAQN	27
consensus		.....	

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 non conserved  
 similar  
  $\geq 50\%$  conserved