

logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 169700 AXE82 RS06025 WP 062332554.1
NTDB id 1047 HON27 RS03110 WP 000389061.1
consensus

MRLCPRFSLDLLVKVKKLVNFVTLGFSCLSPTHAWRSPQSFTESNTLIMFRFYRKQKFKRLQNTLMAAFLALSIIPLTIT
MRLCPRFSLDLLVKVKKLVNFVTLGFSCLSPTHAWRSPQSFTESNTLIMFRFYRKQKFKRLQNTLMAAFLALSIIPLTIT
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80
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logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 169700 AXE82 RS06025 WP 062332554.1
NTDB id 1047 HON27 RS03110 WP 000389061.1
consensus

ALFFLHSHSKDLEQQSTSYLVSVRDNKQQQVIDYMMAKESEVMGFVRSELAYASGGRFYGLVNAFQRLDVSIEAAREHAQ
ALFFLHSHSKDLEQQSTSYLVSVRDNKQQQVIDYMMAKESEVMGFVRSELAYASGGRFYGLVNAFQRLDVSIEAAREHAQ
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160
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logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 169700 AXE82 RS06025 WP 062332554.1
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consensus

QRYITGSGDQIKTSVLPQSSSYVGSERYRLHLKRYHWAYLELLKRSDFDDILLVDIDGNVWYSIYKYDNFGTNLLTGKYQ
QRYITGSGDQIKTSVLPQSSSYVGSERYRLHLKRYHWAYLELLKRSDFDDILLVDIDGNVWYSIYKYDNFGTNLLTGKYQ
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240
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logo

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NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 169700 AXE82 RS06025 WP 062332554.1
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consensus

DTNLGHTFKRLEQTVNEQRKTNEDFTPWVISDFVQEDGKQYAWLGAPIIQQGYLHSYAMFRLPSNAITKLI AEGSSNPSM
DTNLGHTFKRLEQTVNEQRKTNEDFTPWVISDFVQEDGKQYAWLGAPIIQQGYLHSYAMFRLPSNAITKLI AEGSSNPSM
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320
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logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1
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NTDB id 169700 AXE82 RS06025 WP 062332554.1
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consensus

QTI LVGQDHR SRLTSAEIAVEKSKAVVDLALSGKRAVGTYTNTDGEQIIAAYAPINLKNIH WALVVELPEKEAFARVRQ
QTI LVGQDHR SRLTSAEIAVEKSKAVVDLALSGKRAVGTYTNTDGEQIIAAYAPINLKNIH WALVVELPEKEAFARVRQ
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400
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logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1
NTDB id 374 SMU RS06885 WP 002262930.1
NTDB id 169700 AXE82 RS06025 WP 062332554.1
NTDB id 1047 HON27 RS03110 WP 000389061.1
consensus

LEKLFVFAMLT AIVLVVIA SHYLSNFITSPLLKLTWAAERVSAGDLDEAMINTERKDEIGRLAVSFERMQRSIREKISLI
LEKLFVFAMLT AIVLVVIA SHYLSNFITSPLLKLTWAAERVSAGDLDEAMINTERKDEIGRLAVSFERMQRSIREKISLI
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480
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0

logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1	KSQNKELESNLLIIRKQNDQLANKLKDEFLATTSHLRTPLHGMIGIAEALISGANGPISAAHKYQLDIISSGQRLA	560
NTDB id 374 SMU RS06885 WP 002262930.1	KSQNKELESNLLIIRKQNDQLANKLKDEFLATTSHLRTPLHGMIGIAEALISGANGPISAAHKYQLDIISSGQRLA	0
NTDB id 169700 AXE82 RS06025 WP 062332554.1		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
consensus		0

logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1	TLVDDLLDYHKMRYGALDIQRCAYDLSSATRLVLELSHHLLGKKTLRINIQVSEQPVVWSADPQRLEQVLYNLIGNAIKY	640
NTDB id 374 SMU RS06885 WP 002262930.1	TLVDDLLDYHKMRYGALDIQRCAYDLSSATRLVLELSHHLLGKKTLRINIQVSEQPVVWSADPQRLEQVLYNLIGNAIKY	0
NTDB id 169700 AXE82 RS06025 WP 062332554.1		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
consensus		0

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NTDB id 1143 A1552VC RS01965 WP 001883436.1	TSEGKIVISATYIDDKVRVQVVDTGQGI PAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLYVSSQPMV	720
NTDB id 374 SMU RS06885 WP 002262930.1	TSEGKIVISATYIDDKVRVQVVDTGQGI PAEQLEHIFEPLIQAGHDASRYRQGAGLGLSISRQLIELMHGTLYVSSQPMV	0
NTDB id 169700 AXE82 RS06025 WP 062332554.1		0
NTDB id 1047 HON27 RS03110 WP 000389061.1		0
consensus		0

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NTDB id 1143 A1552VC RS01965 WP 001883436.1	GTTFSFTLPLASEEEEIAQARLTLPFHQAPEVLDSELPEQSNLEAAEEMKLTLLYAEPPNLDYLLKSNRRACDTHIF	800
NTDB id 374 SMU RS06885 WP 002262930.1		32
NTDB id 169700 AXE82 RS06025 WP 062332554.1		38
NTDB id 1047 HON27 RS03110 WP 000389061.1		38
consensus		

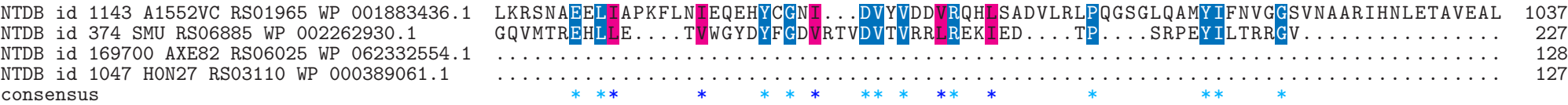
logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1	DGFEALSKIAEENPDLYEYDIMPRLDGYQVCSLRNITXDANYPYIMLSSKDGLEDKVRGREYGSDEYLTKEPFSKELL	880
NTDB id 374 SMU RS06885 WP 002262930.1	DGHQVLEAIKREKPELVLDIMMPGMSGYQVCEKLRQTYDHAELPIIMLTALNQSDDRVRGFEGANDYLSKPFNKQELA	109
NTDB id 169700 AXE82 RS06025 WP 062332554.1	DGREALSKYEEENPDLIILDLMLPELDGLEVAKEVRK...NSHVPIIMLSAKDSEFDKVGLEIGADDYVTKPFSNRELL	118
NTDB id 1047 HON27 RS03110 WP 000389061.1	DGFDALAKIADSNPDLIFVDIMMPRLDGYQTCSLIKNNADFAAKPVIMLSSKDGLFDKARGRIVGSDYELTKPFSKEELF	118
consensus		

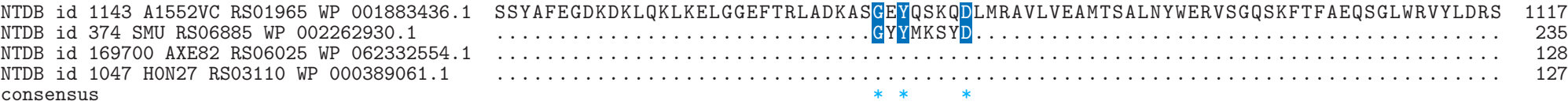
logo

NTDB id 1143 A1552VC RS01965 WP 001883436.1	ARIVAHILTASNAVERIRPRTEAEERAAEANLQKELKHRAMVEASLLETQGRLLIEQLESAPAILCVKEGKVRVFANEAAARLFRRTPEQ	960
NTDB id 374 SMU RS06885 WP 002262930.1	ARIVAHILTASNAVERIRPRTEAEERAAEANLQKELKHRAMVEASLLETQGRLLIEQLESAPAILCVKEGKVRVFANEAAARLFRRTPEQ	175
NTDB id 169700 AXE82 RS06025 WP 062332554.1	ARVKAHLRRTENIESAVAEEN.....ASGIPEIITIGDLQILPDFAVAKKRGTEVELTHREFELLHHLATHT	128
NTDB id 1047 HON27 RS03110 WP 000389061.1	DAIERHRPASNAIRNHVSS.....	127
consensus		

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