

logo

NTDB id 998048 ABDX87 RS18180 WP 346829129.1	MAKAKRMVGGTECGATFPKWAGQCGECGAWNTLVETVIEGGAAAPSGRACWTGQQAQIKTLAEVSVVEETPRFSTNSAEL	80
NTDB id 125 BSU 00870 NP 387968.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	80
NTDB id 537 SMSK321 RS07120 WP 080550752.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
NTDB id 509 SM12261 RS00130 WP 078228442.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
NTDB id 279 KZH43 RS00140 WP 074017595.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
NTDB id 238 SPD RS00140 WP 074017595.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
NTDB id 204 SPR RS00140 WP 074017595.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
NTDB id 169 SP RS00155 WP 074017595.1	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	77
consensus	MAKTKSKFTICQSCGYESPKNMCKCPGCGAWNTMVEEMIKKAPANRRRAAFSHSVQTVQKPSPTSEISEEPRVKTLGEEF	

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NTDB id 998048 ABDX87 RS18180 WP 346829129.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	159
NTDB id 125 BSU 00870 NP 387968.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	160
NTDB id 537 SMSK321 RS07120 WP 080550752.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
NTDB id 509 SM12261 RS00130 WP 078228442.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
NTDB id 279 KZH43 RS00140 WP 074017595.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
NTDB id 238 SPD RS00140 WP 074017595.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
NTDB id 204 SPR RS00140 WP 074017595.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
NTDB id 169 SP RS00155 WP 074017595.1	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	156
consensus	NRVLGGGVVPGSLVLIGGDPGIGKSTLLQVSTQLSQVGTVLVVSIGEESAAQIKLRAERLGDIDSEFYLYAETNMQSVR	

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NTDB id 998048 ABDX87 RS18180 WP 346829129.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	239
NTDB id 125 BSU 00870 NP 387968.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	240
NTDB id 537 SMSK321 RS07120 WP 080550752.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
NTDB id 509 SM12261 RS00130 WP 078228442.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
NTDB id 279 KZH43 RS00140 WP 074017595.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
NTDB id 238 SPD RS00140 WP 074017595.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
NTDB id 204 SPR RS00140 WP 074017595.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
NTDB id 169 SP RS00155 WP 074017595.1	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	236
consensus	AEVERIQPDFLIIDSIQTIMSPEISGVQGSVSQVREVTAEMLQAKTNNIAIFIVGHVTKEGTLAGPRMLEHMDTVLYF	

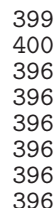
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NTDB id 998048 ABDX87 RS18180 WP 346829129.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	319
NTDB id 125 BSU 00870 NP 387968.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	320
NTDB id 537 SMSK321 RS07120 WP 080550752.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
NTDB id 509 SM12261 RS00130 WP 078228442.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
NTDB id 279 KZH43 RS00140 WP 074017595.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
NTDB id 238 SPD RS00140 WP 074017595.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
NTDB id 204 SPR RS00140 WP 074017595.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
NTDB id 169 SP RS00155 WP 074017595.1	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	316
consensus	EGESDGRILRLRAVKNRFGAVNELGVFGMTDKGLKEVSNPSAIFLTRAQEA VAGSVVMATWEGTRPILAEVQALVPTMF	

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NTDB id 238 SPD RS00140 WP 074017595.1
NTDB id 204 SPR RS00140 WP 074017595.1
NTDB id 169 SP RS00155 WP 074017595.1
consensus

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NTDB id 279 KZH43 RS00140 WP 074017595.1
NTDB id 238 SPD RS00140 WP 074017595.1
NTDB id 204 SPR RS00140 WP 074017595.1
NTDB id 169 SP RS00155 WP 074017595.1
consensus

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- X non conserved
- X similar
- X $\geq 50\%$ conserved