

logo

NTDB id 57206 SAI2T2 RS04620 WP 000959282.1  
NTDB id 26 SA RS04860 WP 000959282.1  
NTDB id 42 MW RS04740 WP 000959293.1  
consensus

MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV  
MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV  
MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGFVKV  
!!\*!!!!

80  
80  
80

logo

NTDB id 57206 SAI2T2 RS04620 WP 000959282.1  
NTDB id 26 SA RS04860 WP 000959282.1  
NTDB id 42 MW RS04740 WP 000959293.1  
consensus

EIEPFLKEIKQIPDILINNKYVIELQYSPIPYKQILQRTEGLKMKGYKVSWLLNDVDYCHNKVKFNHFHSLFINPITRKL  
EIEPFLKEIKQIPDILINNKYVIELQYSPIPYKQILQRTEGLKMKGYKVSWLLNDVDYCHNKVKFNHFHSLFINPITRKL  
EIEPFLKEIKQIPDILINNKYVIELQYSPIPYKQILQRTEGLKMKGYKVSWLLNDVDYCHNKVKFNHFHSLFINPITRKL  
!!\*!\*!\*!\*!!!!

160  
160  
160

logo

NTDB id 57206 SAI2T2 RS04620 WP 000959282.1  
NTDB id 26 SA RS04860 WP 000959282.1  
NTDB id 42 MW RS04740 WP 000959293.1  
consensus

HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIS<sub>S</sub>ELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ  
HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIIELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ  
HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIS<sub>S</sub>ELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ  
!!\*!!

240  
240  
240

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NTDB id 26 SA RS04860 WP 000959282.1  
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consensus

LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIVALESKTAIIEKLINNYLNICSDR  
LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIVALESKTAIIEKLINNYLNICSDR  
LTDQEVVHNYGYIFPEQIYIK<sub>K</sub>NHPIEWQLQVDLWLKNGKSKLVNDNLNYFKLKKFIV<sub>G</sub>LESKTAIIEKLINNYLNICSDR  
!!!!!!!!!!!!!!!!\*!!!!!!!!!!!!!!!!\*!!!!!!!!!!!!!!!!\*!!

320  
320  
320

logo

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NTDB id 26 SA RS04860 WP 000959282.1  
NTDB id 42 MW RS04740 WP 000959293.1  
consensus

GNDVQILF  
GNDVQILF 328  
GNDVQILF 328  
GNDVQILF 328  
!!!!!!!!!!!!

non conserved  
 similar  
 ≥ 50% conserved