

logo

NTDB id 57292 SAI8T7 RS04625 WP 000959282.1
NTDB id 26 SA RS04860 WP 000959282.1
NTDB id 42 MW RS04740 WP 000959293.1
consensus

MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV
MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV
MLVALNEEKERVLATTALRKTQYFCPVCGKQVILKRGLKVIISHFAHKHLAEQKCFNNETIKHYKSKLILAQMIQQQGCKV
!!*!!!!

80
80
80

logo

NTDB id 57292 SAI8T7 RS04625 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 57292 SAI8T7 RS04625 WP 000959282.1
NTDB id 26 SA RS04860 WP 000959282.1
NTDB id 42 MW RS04740 WP 000959293.1
consensus

HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIS_SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ
HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIS_SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ
HTFNLEKKQIMMFQQIQYLGGHKYVAEKRNAKIS_SELFNEAPCDYHAVYKLSKFAINQYIKYCRWQNSVLEPTLSAMYQLQ
!!*!!

240
240
240

logo

NTDB id 57292 SAI8T7 RS04625 WP 000959282.1
NTDB id 26 SA RS04860 WP 000959282.1
NTDB id 42 MW RS04740 WP 000959293.1
consensus

LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLNKNGKSKLVNDNLNYFKLKKFIVALESKTAIIEKLINNYLNICSDR
LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLNKNGKSKLVNDNLNYFKLKKFIVALESKTAIIEKLINNYLNICSDR
LTDQEVVNYGYIFPEQIYIENHPIEWQLQVDLWLNKNGKSKLVNDNLNYFKLKKFIVALESKTAIIEKLINNYLNICSDR
!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!*!!!!!!!!!!!!!!!!*!!

320
320
320

logo

NTDB id 57292 SAI8T7 RS04625 WP 000959282.1
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