logo
NTDB id 352764 E3983 RS04215 WP 245183405.1 MLQIFASITQRLRFPTVCLLCKQYQRDAQFICKECLCLFKPIKYGCKYCACPLPDAHFPVCGACSTKQPDFDRAIIAYPY 80
NTDB id 1270 LPP RS11540 WP 015961431.1 MLQKFLSITQNLRLPSICTLCNQFHKSQLAVCSNCMEFMKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF 80
NTDB id 1247 GCO85 RS11680 WP 015961431.1 .....MKQLGSCCQYCAYPLSDDTYLVCGQCVRKRPSFDSAYIAYRF 42
consensus
logo
NTDB id 352764 E3983 RS04215 WP 245183405.1 EEPLRTLIHQFKYQEKLYLVSFLSQLMLPFLKECDEIPQCLIPVPMHPQRLKQRGFNQAAVLTKSLAKKMHIPYQLQGCQ 160
NTDB id 1270 LPP RS11540 WP 015961431.1 EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ 160
NTDB id 1247 GCO85 RS11680 WP 015961431.1 EEPLRSLIHQFKYHNGLYLASFLKQLLLNALPKSALKPDCLIPVPMHPKRLKRRGFNQAAVLTRLLARQLNIPYDLYYCQ 122
consensus
logo
NTDB id 352764 E3983 RS04215 WP 245183405.1 KIINTKPQASLDKKQRAKNLRQAFTVNTLPYQHIALVDDLLTTGNTANELAQAFKKKGVLRVDVWCCARAVLA. 233
NTDB id 1270 LPP RS11540 WP 015961431.1 KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN 234
NTDB id 1247 GCO85 RS11680 WP 015961431.1 KIINTASQANLDGEQRRKNLRHAFYVPPVTYEHVMIVDDLLTTGSTANEIAHTLKNAGVKRVDICCCARAVTKN 196
consensus
X non conserved
X similar
X ≥ 50% conserved