

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

MLASKMRWEIQRPDQDKVKSLTEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQ

80
80

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

[illegible]

60
60

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

GLDVIITDHHHEPGPELPDVRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI
GLDVIITDHHHEPGPELPDVRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI

240
240

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

[illegible]20
20

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

RQK**M**VSKMTDEAIDMVEQQGLDQTAI**V**VAKAGWNPVGIVASKLVDRFYRPAI**V**LGIDEEEKGI**A**KGSARSIRGFNLFES
RQKMVSKMTDEAIDMVEQQGLDQTAI**V**VAKAGWNPVGIVASKLVDRFYRPAI**V**LGIDEEEKGI**A**KGSARSIRGFNLFES
RQKMVSKMTDEAIDMVEQQGLDQTAI**V**VAKAGWNPVGIVASKLVDRFYRPAI**V**LGIDEEEKGI**A**KGSARSIRGFNLFES
!!!!!!!!!!!!!!*

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NTDB id 354 BSU 27620 NP 390640.1
consensus

LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN

:80
:80

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGLEEGIVPGSRISIVGEMSIENEWNNRKKPQQLMIKDA
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGLEEGIVPGSRISIVGEMSIENEWNNRKKPQQLMIKDA

60
60

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



640
640

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



720
720

logo

NTDB id 981488 NSU09 RS16275 WP 029727133.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



786
786

 non conserved
 similar
 ≥ 50% conserved