

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ

80
80

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

60
60

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

GLDVIITDHHHEPGPELPNVRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI

240
240

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE

20
20

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consensus

LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
 LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN
 LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVLDVCGVEDITVESIAEMNMLSPFGMLN

:80
:80

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus

[illegible]60
60

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



640
640

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



720
720

logo

NTDB id 340575 ETK61 RS14850 WP 069837662.1
NTDB id 354 BSU 27620 NP 390640.1
consensus



786
786

X non conserved
X similar
X ≥ 50% conserved