

logo

NTDB id 340824 ETL58 RS18475 WP 101172232.1
NTDB id 108 BSU 35470 NP 391427.1
consensus

MNVPEKNSSFSKELQQRLRSRHLRLRTELFSFSDMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSGKN
MNVPEKNSSFSKELQQRLRSRHLRLRTELFSFSDMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSGKN
MNVPEKNSSFSKELQQTLRSRHLRLRTELFSFSDMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSGKN
!!

80
80

logo

NTDB id 340824 ETL58 RS18475 WP 101172232.1
NTDB id 108 BSU 35470 NP 391427.1
consensus

KLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
KLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
KLYCRSCVMMGRVSEEVPLYSWKEENESNWKSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
!!

160
160

logo

NTDB id 340824 ETL58 RS18475 WP 101172232.1
NTDB id 108 BSU 35470 NP 391427.1
consensus

ESALNQLRVCIA TPRTDWLELAPRLKAAFQGAESALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
ESALNQLRVCIA TPRTDVVLELAPRLKAAFQGAESALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
ESALNQLRVCIA TPRTDVVLELAPRLKAAFQGAESALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
!!

240
240

logo

NTDB id 340824 ETL58 RS18475 WP 101172232.1
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consensus

SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
!!

320
320

logo

NTDB id 340824 ETL58 RS18475 WP 101172232.1
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consensus

WIKFHVKEGRPVFLFVPSVSI LEKAAACFRGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
WIKFHVKEGRPVFLFVPSVSI LEKAAACFRGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
WIEFHVKEGRPVFLFVPSVSI LEKAAACFRGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
!!

400
400

logo

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consensus

LGAESPIFTESALVQIAGRTGRHKEYADGDVIYFHFHGKTKSMLDARKHIKEMNELAAKVECTD
LGAESPIFTESALVQIAGRTGRHKEYADGDVIYFHFHGKTKSMLDARKHIKEMNELAAKVECTD
LGAESPIFTESALVQIAGRTGRHKEYADGDVIYFHFHGKTKSMLDARKHIKEMNELAAKVECTD
!!

463
463

⊠ non conserved
⊠ similar
⊠ ≥ 50% conserved