

logo

NTDB id 336406 EQY76 RS18470 WP 003243962.1
NTDB id 108 BSU 35470 NP 391427.1
consensus

MNVPEKNSSFSKELQQTLLRSRHLRLTELFSFSDMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSGKN
MNVPEKNSSFSKELQQTLLRSRHLRLTELFSFSDMIEWHIKNGYITAENSISINKRRYRCNRCGQTDQRYFSFYHSSGKN
!!

80
80

logo

NTDB id 336406 EQY76 RS18470 WP 003243962.1
NTDB id 108 BSU 35470 NP 391427.1
consensus

KLYCRSCVMMGRVSEEVPLYSWKEENESNWSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
KLYCRSCVMMGRVSEEVPLYSWKEENESNWSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
KLYCRSCVMMGRVSEEVPLYSWKEENESNWSIKLTWDGKLSGGQQAANVLEAISKKEELLIWAVCGAGKTEMLFPGI
!!

160
160

logo

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NTDB id 108 BSU 35470 NP 391427.1
consensus

ESALNQLRVCIA TPRTDWLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
ESALNQLRVCIA TPRTDWLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
ESALNQLRVCIA TPRTDWLELAPRLKAAFQGADISALYGGSDDKGRLSPLMISTTHQLLRYKDAIDVMIIDEVDAFPY
!!

240
240

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consensus

SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
SADQTLQFAVQKARKKNSTLVYLSATPPKELKRKALNGQLHSVRI PARHHRKPLPEPRFVWCGNWKKKLNRNKIPPAVKR
!!

320
320

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consensus

WIEFHVKEGRPVFLVPSVSI LEKAAACFKGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
WIEFHVKEGRPVFLVPSVSI LEKAAACFKGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
WIEFHVKEGRPVFLVPSVSI LEKAAACFKGVHCR TASVHAEDKHRKEKVQQFRDGLDLLITTTIL ERGVTVPKVQTGV
!!

400
400

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consensus

LGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD
LGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD
LGAESSIFTESALVQIAGRTGRHKEYADGDVIYFHFGKTKSMLDARKHIKEMNELAAKVECTD
!!

463
463

non conserved
 similar
 ≥ 50% conserved