

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDADFYDPPFEMKGMKEAADRIKQAISSQ  
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDADFYDPPFEMKGMKEAADRIKQAISSQ  
!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

80  
80

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL  
EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL  
EKIMYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIKERGFSLIITVDTGIAAVHEAKVAKEL  
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160  
160

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

GLDVIITDHHHEPGPELPDVRRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI  
GLDVIITDHHHEPGPELPDVRRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI  
GLDVIITDHHHEPGPELPDVRRAIVHPKQPGCTYPFKELAGVGVAFLAHALLGELPDELLDLAAIGTIADLVPLHDENRLI  
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240  
240

logo

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NTDB id 354 BSU 27620 NP 390640.1  
consensus

ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE  
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE  
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE  
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320  
320

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

RQKMVSKMTDEAIEMVQQGLDQTAIVVAKAGWNPGVVGIVASKLVDRFYRPAIVLGIDEEKGIKGSARSIRGFNLFES  
RQKMVSKMTDEAIEMVQQGLDQTAIVVAKAGWNPGVVGIVASKLVDRFYRPAIVLGIDEEKGIKGSARSIRGFNLFES  
RQKMVSKMTDEAIEMVQQGLDQTAIVVAKAGWNPGVVGIVASKLVDRFYRPAIVLGIDEEKGIKGSARSIRGFNLFES  
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400  
400

logo

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consensus

LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVDLVCGVEDITVESIAEMNMLSPFGMLN  
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVDLVCGVEDITVESIAEMNMLSPFGMLN  
LSECRDILPHFGGHPMAAGMTLKAEDVPDLRSRLNEIADNTLTEEDFIPVQEVDLVCGVEDITVESIAEMNMLSPFGMLN  
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480  
480

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
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consensus

PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNRKKPQLMIKDA  
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNRKKPQLMIKDA  
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENWNRKKPQLMIKDA  
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560  
560

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus



640  
640

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus



720  
720

logo

NTDB id 87135 BsBEST3145 RS14730 WP 003246110.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus



786  
786

 non conserved  
 similar  
 ≥ 50% conserved