

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ  
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ  
MLASKMRWEIQRPDQDKVKSLEQLHITPLVASLLVKGFDTAESARLFLHTKDAFDYDPFEMKGMKEAADRIKQAISSQQ

80  
80

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

EKIMIIYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIK**D**RGFSLIITVDTGIAAVHEAKVAKEL  
 EKIMIIYGDYDADGVTSTSVMLHTLQKLSAQVDFYIPDRFKEGYGPNEQAFRSIK**D**RGFSLIITVDTGIAAVHEAKVAKEL  
 !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!\*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

60  
60

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

[illegible]240  
240

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
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consensus

ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE  
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE  
ATLGLERLRRTNRLGLKELIKLSGGDIGEANEETVGFQLAPRLNAVGRIEQADPAVHLLMSEDSFEEELAAEIDQLNKE

20  
20

logo

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consensus

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NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

80  
80

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus

PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENEWNNRKKPQLMIKDA  
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENEWNNRKKPQLMIKDA  
PKPHVLVENAVLEDVRKIGANKTHVKMTIRNESSQLDCVGFNKGELQEGIVPGSRISIVGEMSIENEWNNRKKPQLMIKDA

60  
60

logo

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consensus



640  
640

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus



720  
720

logo

NTDB id 319605 D9C16 RS21795 WP 014480432.1  
NTDB id 354 BSU 27620 NP 390640.1  
consensus



786  
786

 non conserved  
 similar  
 ≥ 50% conserved