

logo

NTDB id 1148353 LSAF01 RS04970 WP 011374694.1  
NTDB id 626 LCA RS04960 WP 011374694.1  
consensus

MLTKNQLLLKVHLAGFGTISELRLAAWLSTSHNWALSALEIAQIARLPERYWPTFQASFQSTTLQRQCIDHEVRTTYLT  
MLTKNQLLLKVHLAGFGTISELRLAAWLSTSHNWALSALEIAQIARLPERYWPTFQASFQSTTLQRQCIDHEVRTTYLT  
MLTKNQLLLKVHLAGFGTISELRLAAWLSTSHNWALSALEIAQIARLPERYWPTFQASFQSTTLQRQCIDHEVRTTYLT  
!!

80  
80

logo

NTDB id 1148353 LSAF01 RS04970 WP 011374694.1  
NTDB id 626 LCA RS04960 WP 011374694.1  
consensus

ILDKDYPQRLLLETYLPPVLLFYRGDLRLLKQPCLAVVGARQATHYSKQSLEQLLQGLTATTIISGLAQGADAMAHEVALQ  
ILDKDYPQRLLLETYLPPVLLFYRGDLRLLKQPCLAVVGARQATHYSKQSLEQLLQGLTATTIISGLAQGADAMAHEVALQ  
ILDKDYPQRLLLETYLPPVLLFYRGDLRLLKQPCLAVVGARQATHYSKQSLEQLLQGLTATTIISGLAQGADAMAHEVALQ  
!!

160  
160

logo

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consensus

RGLAPIGVIGTGINCSYPPQNEHLQQVAEKGLLSEYALGTPARRYQFPARNKIIAGLCHSLIVTEARHKSGSLITANL  
RGLAPIGVIGTGINCSYPPQNEHLQQVAEKGLLSEYALGTPARRYQFPARNKIIAGLCHSLIVTEARHKSGSLITANL  
RGLAPIGVIGTGINCSYPPQNEHLQQVAEKGLLSEYALGTPARRYQFPARNKIIAGLCHSLIVTEARHKSGSLITANL  
!!

240  
240

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consensus

ALQANRNWYAIPGRIDQSLSQGCNQLIAAGATPLLDKNILIEELRYFD  
ALQANRNWYAIPGRIDQSLSQGCNQLIAAGATPLLDKNILIEELRYFD  
ALQANRNWYAIPGRIDQSLSQGCNQLIAAGATPLLDKNILIEELRYFD  
!!

288  
288

- non conserved
- similar
- ≥ 50% conserved