

85

VQRP^gSLHⁱRR^qLS^lLN^rAS^sIS^tLS^hFS^fCL^qVL^kK^kY^yYL^lIDL^dDP^pGF^rIA^dQ^qTE^gEL^lIG^dEV^lDEL^fFE^dEY^aK^gEKA^fFF^eLV^dRY^tTT

170

DRHDLDLQFLVKQVVEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRALELTKAPGGPAPRAD
DRHDLDLQFLVKQVVEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRALELTKAPGGPAPRAD
DRHDLDLQFLVKQVVEYSRSHPNPEAWLESFVHLYDVSEKSAIEELPFYQYVKEDIAMVLNGAKEKLLRALELTKAPGGPAPRAD

255

NFLDDLAQIDELIQHDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLEKLKTDYFTRSPEQHLKSLAEMKP
NFLDDLAQIDELIQHDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLEKLKTDYFTRSPEQHLKSLAEMKP
NFLDDLAQIDELIQHDDFSELYKRVPAVSFKRAKAVKGDEFDPALLDEATDLRNGAKKLEKLKTDYFTRSPEQHLKSLAEMKP

340

VIETLVQLVISYGKRFEAAKQEKSIIDFSDLEHYCLAILTAENDKGEREPSEAAIFYQQFHEVLVDEYQDTNLVQESILQLVTS
VIETLVQLVISYGKRFEAAKQEKSIIDFSDLEHYCLAILTAENDKGEREPSEAAIFYQQFHEVLVDEYQDTNLVQESILQLVTS
VIETLVQLVISYGKRFEAAKQEKSIIDFSDLEHYCLAILTAENDKGEREPSEAAIFYQQFHEVLVDEYQDTNLVQESILQLVTS

425

GPEETGNLFMVGDVKQSIYRFRLAEP¹LLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSTNFLFKQLMGGKIGVDYDEQAE
 GPEETGNLFMVGDVKQSIYRFRLAEP¹LLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSTNFLFKQLMGGKIGVDYDEQAE
 GPEETGNLFMVGDVKQSIYRFRLAEP¹LLFLSKYKRFTESGEGTGRKIDLNKNFRSRADILDSTNFLFKQLMGGKIGVDYDEQAE

510

LKLGAAYPDNDDETETELLIDNAEDTDASEEAAELETQFEAKAIAKEIRKLSSPFKVYDGKKKTHRNQYRDIVILLRSMPTWALKLGAAYPDNDDETETELLIDNAEDTDASEEAAELETQFEAKAIAKEIRKLSSPFKVYDGKKKTHRNQYRDIVILLRSMPTWALKLGAAYPDNDDETETELLIDNAEDTDASEEAAELETQFEAKAIAKEIRKLSSPFKVYDGKKKTHRNQYRDIVILLRSMPTWAL

595



logo

NTDB id 53648 K6K01 RS05740 WP 003233100.1
NTDB id 119 BSU 10630 NP 388944.2
consensus



1232
1232

 non conserved
 similar
 ≥ 50% conserved