logo

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NTDB id 521 SMSK321 RS04960 WP 000443804.1   MESIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPHTPVKETNLQAEVVAISKDSSTEKDVKKEEKEEPVEQDLITVDVKG   80
NTDB id 493 SM12261 RS04300 WP 000387351.1   MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVAVSKDSVSEKEVKKEEKEEPLEQDLITVDVKG   80
NTDB id 153 SP RS04730 WP 000387344.1        MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVAAVSKDSSTEKEVKKEEKEEPLEQDLITVDVKG   80
NTDB id 263 KZH43 RS04195 WP 000387330.1     MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVAAVSKDLVSEKEVNKEEKEEPLEQDLITVDVKG   80
NTDB id 222 SPD RS04535 WP 000387330.1       MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVAAVSKDLVSEKEVNKEEKEEPLEQDLITVDVKG   80
NTDB id 188 SPR RS04295 WP 000387330.1       MEAIIEKIKEYKIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVAAVSKDLVSEKEVNKEEKEEPLEQDLITVDVKG   80
NTDB id 183182 LLUC11 RS09025 WP 237025748.1 MDKILEKVKEYWKMIVLVVCGLIAGGIFYVLTNGQKPTTNLSVENL.SSVSRQSSV...SKFSEP.NEKSVSKIMVDLKV   75
NTDB id 595 KW2 RS08725 WP 021037687.1       MEKILDKLKEYWKIMLLVVCALIAGGIFYIFTNSPKPAETLSVENL.SSSSTKSSV...SKFNSSSSEKNKNEIMVDLKG   76
consensus                                    !**!**!*!!!**********!**!!*!********!********** **!*** **********************!*!!*!*
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NTDB id 521 SMSK321 RS04960 WP 000443804.1   AVKSPGIYDLSVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEESASQHTGSGAPSST......SKDKK   154
NTDB id 493 SM12261 RS04300 WP 000387351.1   AVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAASQQAGPGTASST......SKEKK   154
NTDB id 153 SP RS04730 WP 000387344.1        AVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGSGTASST......SKEKK   154
NTDB id 263 KZH43 RS04195 WP 000387330.1     AVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGLGTASSI......SKEKK   154
NTDB id 222 SPD RS04535 WP 000387330.1       AVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGLGTASSI......SKEKK   154
NTDB id 188 SPR RS04295 WP 000387330.1       AVKSPGIYDLPVGSRVNDAVQKAGGLTEQADSKSLNLAQKVSDEALVYVPTKGEEAVSQQTGLGTASSI......SKEKK   154
NTDB id 183182 LLUC11 RS09025 WP 237025748.1 AVTKPNVYQISSDERLVDLIKEAGGFTDQADQKSINLSAKLKDEEVIYVPKIGENSSTENTDSSANSVTS..........   145
NTDB id 595 KW2 RS08725 WP 021037687.1       AVAKPNVYQISSDERLVDLIRQAGGFTDQADQKSINLSAKLKDEEVIYVPKVGESSSSESTDSPTGSSVSNQISTTSGPK   156
consensus                                    !!**!**!******!**!****!!!*!*!!!*!!*!!**!**!!***!!!**!!** ***** ***!*       *****
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logo

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NTDB id 521 SMSK321 RS04960 WP 000443804.1   INLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGAKTIEKLKDYVTVD   216
NTDB id 493 SM12261 RS04300 WP 000387351.1   VNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGAKTIEKLKDYVTVD   216
NTDB id 153 SP RS04730 WP 000387344.1        VNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 263 KZH43 RS04195 WP 000387330.1     VNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 222 SPD RS04535 WP 000387330.1       VNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 188 SPR RS04295 WP 000387330.1       VNLNKASLEELKQVKGLGGKRAQDIIDHREANGKFKSVDELKKVSGIGGKTIEKLKDYVTVD   216
NTDB id 183182 LLUC11 RS09025 WP 237025748.1 ............................................................   145
NTDB id 595 KW2 RS08725 WP 021037687.1       VNINKADLTELQKLTGIGQKKAQDIIDFRMKNGDFKSIEDLGKVSGFGDKTLEKLKDEISID   218
consensus                                    ************************************************************
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X non conserved

X similar

X ≥ 50% conserved