

NTDB id 613	LCA RS05340 WP 231952379.1	MMWQDYWQRYYRFFWWVPCLVLGLIIAGGFYWHQSQQKQKVITQ.N.EQLLAT	SASKTKP...ATKKTHTQAATKKKGGFVDEKGA	80
NTDB id 521	SMSK321 RS04960 WP 000443804.1	..MESIIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPHTPVKETNLQAEVVA	ISKDSSSTEKDVKKKEEKKEEPVEQDLITVDVKGA	83
NTDB id 493	SM12261 RS04300 WP 000387351.1	..MEAIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVA	AVSKDSVSEKEVKKKEEKKEEPLEQDLITVDVKGA	83
NTDB id 153	SP RS04730 WP 000387344.1	..MEAIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVA	AVSKDSSSTEKEVKKKEEKKEEPLEQDLITVDVKGA	83
NTDB id 263	KZH43 RS04195 WP 000387330.1	..MEAIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVA	AVSKDLVSEKEVNKEEKKEEPLEQDLITVDVKGA	83
NTDB id 222	SPD RS04535 WP 000387330.1	..MEAIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVA	AVSKDLVSEKEVNKEEKKEEPLEQDLITVDVKGA	83
NTDB id 188	SPR RS04295 WP 000387330.1	..MEAIIEKIKIKEYKIIIVICTGLGLLVGGFFLLKPAPQTPVKETNLQAEVVA	AVSKDLVSEKEVNKEEKKEEPLEQDLITVDVKGA	83
NTDB id 20412	L RS09240 WP 010906131.1	..MDKILEKVKEYWKMIIVLVVCGLIAGGIFYVLTNGQKPTTNLSVENL.SSVSRQSSV...	SKFSEPN.EKSVSKIMVDLKGA	78
NTDB id 595	KW2 RS08725 WP 021037687.1	..MEKILDKLKEYWKIMLLVVCALIIAGGIFYIFTNSPKPAETLSVENL.SSSSTKSSV...	SKFNSSSSEKKNBIMVDLKGA	79
consensus		*****	*****	

[illegible][illegible]

-  non conserved
-  similar
-  >50% conserved