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NTDB id 615 LCA RS02545 WP 011374200.1	MKVNLDYLGRFLTENELTE	EERQLAEKLPAMRKEKGKLFQRCNSTIL	EEWYLPIGAYYCRECLLMKRV	74
NTDB id 593 KW2 RS05130 WP 021037147.1	MGQQVIACGRQFTAAQLADTQN...	NNYSLPQIKRPAFLRVKHLVQCRCQVVP	PQTCLPDGRHYCAQCLLFGRL	80
NTDB id 115754 SMUFR RS02365 WP 002263026.1	MNLILENLGLHLLLEKDISAFNNFTDQVNN	ENKLIKIGAMTSVNANKVRCNRCGTIHIKTNVKLPIGAFFCPTCLELGRV		69
NTDB id 277 KZH43 RS10090 WP 000867601.1	MENLENYVGRLLTENQMNS	DLKKQAKILPAMIKKPHYFICNRCSQNN	LGNALPDGSIYCRACLVFGR	69
NTDB id 236 SPD RS10765 WP 000867601.1	MKVNLDYLGRFLTENELTE	EERQLAEKLPAMRKEKGKLFQRCNSTIL	EEWYLPIGAYYCRECLLMKRV	69
NTDB id 202 SPR RS10250 WP 000867601.1	MKVNLDYLGRFLTENELTE	EERQLAEKLPAMRKEKGKLFQRCNSTIL	EEWYLPIGAYYCRECLLMKRV	69
NTDB id 167 SP RS11275 WP 000867616.1	MKVNLDYLGRFLTENELTE	EERQLAEKLPAMRKEKGKLFQRCNSTIL	EEWYLPIGAYYCRECLLMKRV	69
NTDB id 507 SM12261 RS09240 WP 000867722.1	MKVNPNYLGRFLTENELTE	EERQLAEKLPAMRKEKGKLFQRCNSTIL	EEWYLPIGAYYCRECLLMKRV	69
NTDB id 535 SMSK321 RS10515 WP 000867726.1	MKVNPNYLGRFLTENELTK	EERQLAEKLPAMRKEKGKLFQRCDSAIL	DEWYLPIGAYYCRECLLMKRV	69
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NTDB id 615 LCA RS02545 WP 011374200.1	VEGDWLYTVPECHLFENNIPKLTWEGQLTPHQEKA	AQAVVEVVQTQKRHV	LTAVTGAGKTEMLFQGITLVALQKQGRVCLA	154
NTDB id 593 KW2 RS05130 WP 021037147.1	RSDYFYHLPQQDFPEK..TYLRWTGKLTENQEKISDALCQQITNNQKLLVQAVTGAGKTEMIYQLIEQILSHGGSVGLA			158
NTDB id 115754 SMUFR RS02365 WP 002263026.1	TNRDSLYYFEQKPFQK..QVLRWQQLTPFQEVSRGLKKSVCHEKNMLITHAVTGAGKTEMIYETLASILNQGGAVALA			147
NTDB id 277 KZH43 RS10090 WP 000867601.1	RSDQILYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
NTDB id 236 SPD RS10765 WP 000867601.1	RSDQILYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
NTDB id 202 SPR RS10250 WP 000867601.1	RSDQILYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
NTDB id 167 SP RS11275 WP 000867616.1	RSDQILYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
NTDB id 507 SM12261 RS09240 WP 000867722.1	RSDQILYYFPQEDFPKQ..DVLKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
NTDB id 535 SMSK321 RS10515 WP 000867726.1	RSDQVLYYFPQEDFAKQ..DILKWRGQLTPFQEKVSEGLLQAVDKQKPTLVHAVTGAGKTEMIYQVVAKVINAGGAVCLA			147
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NTDB id 615 LCA RS02545 WP 011374200.1	SPRIDVCLLEYKRLQDFSCIA	LLHGESEPYFRTPLVATTHQLLKFYQAFDILLVDEVDAPFPYVDNPMLYHAVKNS	234	
NTDB id 593 KW2 RS05130 WP 021037147.1	APRVAVCLELYPRLQAAFATTSIMLMHGEQTEPYRYTQLVICTTHQLLKFYHAFDITVIVDEVDAPFPYVDNPVLATAVEQA		236	
NTDB id 115754 SMUFR RS02365 WP 002263026.1	SPRIDVCLIELHQRLSRDFTCIQIP..LLYHE.GDSYFRSPLVVMTSHQLLRFKEAFDILLIIDEVDAPFPRDNDMLYFALENA		225	
NTDB id 277 KZH43 RS10090 WP 000867601.1	SPRIDVCLIELYKRLSRDFSCPIIS..LLYGE.SEAYERSPLVATTHQLLKFYRAFDLLIIDEVDAPFPYVDNKMLYYAVDHC		225	
NTDB id 236 SPD RS10765 WP 000867601.1	SPRIDVCLLEYKRLQQDFSCGIA..LLHGE.SEPYFRTPLVATTHQLLKFYQAFDILLIIDEVDAPFPYVDNPMLYHAVKNS		225	
NTDB id 202 SPR RS10250 WP 000867601.1	SPRIDVCLLEYKRLQQDFSCGIA..LLHGE.SEPYFRTPLVATTHQLLKFYQAFDILLIIDEVDAPFPYVDNPMLYHAVKNS		225	
NTDB id 167 SP RS11275 WP 000867616.1	SPRIDVCLLEYKRLQQDFSCGIA..LLHGE.SEPYFRTPLVATTHQLLKFYQAFDILLIIDEVDAPFPYVDNPMLYHAVKNS		225	
NTDB id 507 SM12261 RS09240 WP 000867722.1	SPRIDVCLLEYKRLQQDFSCGIA..LLHGE.SEPYFRTPLVATTHQLLKFYQAFDILLIIDEVDAPFPYVDNPMLYHAVKNS		225	
NTDB id 535 SMSK321 RS10515 WP 000867726.1	SPRIDVCLLEYKRLQDFACEIS..LLYGE.SEPYFRTPLVSTTHQLLKFYQAFDILLIIDEVDAPFPYVDNPMLYHAVKNS		225	
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NTDB id 615 LCA RS02545 WP 011374200.1	CKPQCALLYLTATPTPAIKRAIAAKQMTVSELP	PLRFHGHLLPEPQRHVAFNWRAQLKKGRLPKRLERDCQTYLKT.QQIL	313	
NTDB id 593 KW2 RS05130 WP 021037147.1	KKINGTLLYLTATSTDKLDKQIKKHEIKRFL	PPRRFHGHPLVIPMFFWKKTFYK.....KFIEQRKTCFP	303	
NTDB id 115754 SMUFR RS02365 WP 002263026.1	LKSDGVKVFLLTATSTDQLDKQVKKQKGLKLLH	LARRFHANPLVVPKPIWNLNLSLERLQRGKLP	305	
NTDB id 277 KZH43 RS10090 WP 000867601.1	VKENGRLIFLTATSTNELDKKVRIGELKRLN	PPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKLSYIEKQRKTAAYPLL	305	
NTDB id 236 SPD RS10765 WP 000867601.1	VKENGRLIFLTATSTNELDKKVRIGELKRLN	PPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKLSYIEKQRKTAAYPLL	305	
NTDB id 202 SPR RS10250 WP 000867601.1	VKENGRLIFLTATSTNELDKKVRIGELKRLN	PPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKLSYIEKQRKTAAYPLL	305	
NTDB id 167 SP RS11275 WP 000867616.1	VKENGRLIFLTATSTNELDKKVRIGELKRLN	PPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKLSYIEKQRKTAAYPLL	305	
NTDB id 507 SM12261 RS09240 WP 000867722.1	VKENGRLIFLTATSTNELDKKVRIGELKRLS	PPRRFHGNPLIIPKPIWLSDFNRYLDKNRLSPKLSYIEKQRKTCYPLL	305	
NTDB id 535 SMSK321 RS10515 WP 000867726.1	VKENGRLIFLTATSTDELDRKVRIGELKRLS	PPRRFHGNPLIIPKPIWLSDFNRYLEKKCLSPKLSYIEKQRKTCYPLL	305	
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