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NTDB id 1159	ABDM36	RS04765	WP 001166105.1	VEEGIVITVNINTASAEELATLLKGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEAIVR	95
NTDB id 1150	A1552VC	RS08580	WP 001166105.1	VEEGIVITVNINTASAEELATLLKGIGLKKAAQIVDYREANGPFTHIDDLTNVKGIGEAIVR	95
NTDB id 1395	DSB67	RS04445	WP 010649155.1	KYEGEIVTVNNSATAQEIATLLIGIGEKKAAQIVDYREANGPFKTAADLTKVKGIGEAIVR	87
NTDB id 1362	HI 1008	AAC22669.1		EAQVAPAVVSDKLNINTATASEIQKSLTGIGAKKAEIVQYREKHGNFXNAEQLELVQGIGKATLE	104
NTDB id 1382	A4U84	RS07375	WP 062924234.1	VAATQKVNANVNVNINTATASEIQDKLVGIGAKKAAQIVDYREANGKFLNVEQLTEVSGIGKATLE	104
NTDB id 595	KW2	RS08725	WP 021037687.1	SESTDSPITG...SSVSNQISTTSGPKVNINKADITELQK.LTGIGQKKAQIIDFRMKNQDFKSIDLQKVSQFQDKTLE	209
NTDB id 521	SMSK321	RS04960	WP 000443804.1	SQHTGSGAP.....SSTSCKDKLNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGAKTIE	207
NTDB id 493	SM12261	RS04300	WP 000387351.1	SQQAGPGTA.....SSTSKEKKVNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGAKTIE	207
NTDB id 263	KZH43	RS04195	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGGKTIE	207
NTDB id 222	SPD	RS04535	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGGKTIE	207
NTDB id 188	SPR	RS04295	WP 000387330.1	SQQTGLGTA.....SSISKEKKVNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGGKTIE	207
NTDB id 153	SP	RS04730	WP 000387344.1	SQQTGSGTA.....SSTSKEKKVNINKASLEELKQ.VKGLGGKRAQIIDHREANGKFKSVDELKVVSGIGGKTIE	207
NTDB id 1069	ABD1	RS03060	WP 043960841.1	LNISKSSHSNSSKISQTKNFNDSTGQIHLNQAIVNVEFQQ.LKGIGEKKAQIVDYREANGKFKSVDELKVVSGIGGKTIE	128
NTDB id 1041	HON27	RS14740	WP 005124713.1	LNITKSSHSNGSKISQTKNFNDSTGQIHLNQAIVNVEFQQ.LKGIGEKKAQIVDYREANGKFKSVDELKVVSGIGGKTIE	128
NTDB id 36	MW	RS08230	WP 001793467.1	EP...QIGVNS..VHVKNGTNTNTKVNINTASVSELMS.VPGVQAKANAIVDYREANGKFKSVDELKVVSGIGGKTIE	220
NTDB id 20	SA	RS08005	WP 001802022.1	EP...QIEVNS..VHVKNGTNTNTKVNINTASVSELMS.VPGVQAKANAIVDYREANGKFKSVDELKVVSGIGGKTIE	220
NTDB id 105	BSU	25590	NP 390437.1	V...QQGGGS..V..QSDGGKALVNINTATLEELQ.LSGVGPSKAEIAYREANGKFKSVDELKVVSGIGGKTIE	196
NTDB id 1014	ACIAD	RS13855	WP 004924483.1	LNSSASHHLSLK..P..TQQSGTGKVNINSAITAKELQQ.LHGVEKKALAIYREANGKFKSVDELKVVSGIGGKTIE	120
NTDB id 1127	NGFG	RS10560	WP 003703428.1	SCAFSLAAVNINAASQKLEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLA	105
NTDB id 1126	NGFG	RS09375	WP 003703428.1	SCAFSLAAVNINAASQKLEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLA	105
NTDB id 1125	NGFG	RS07215	WP 003703428.1	SCAFSLAAVNINAASQKLEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLA	105
NTDB id 1108	NGFG	RS06525	WP 003703428.1	SCAFSLAAVNINAASQKLEA.LPGIGPAKAKAIAEYRAQNGAFKSVDDLIKVKGIGPAVLA	105
NTDB id 156435	CjjRM3196	RS00050	WP 002855574.1	LTAFLFGAVNINTATLTKELKS.LNGIGEAKAKAILEYRKE.ANFTSIDDLKVKVGIGDKLFE	70
NTDB id 1242	Cj0011c	YP 002343483.1		LTAFLFGAVNINTATLTKELKS.LNGIGEAKAKAILEYRKE.ANFTSIDDLKVKVGIGDKLFE	70

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NTDB id 1242	Cj0011c	YP 002343483.1		KIKNDITIE.....	79

consensus

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 non conserved  
 similar  
  $\geq 50\%$  conserved