

logo

T10122	AZV00182.1	MK291441:c9463-9056	MRYVAF	LHTD	EAGGFGISFPDFPGA	ISDGD	TVEQAIQR	GERALAF	HVQC	GMREDG	DVPA	PRSV	DILAD	PDLA	EWREGA	Q	80				
T1039	NP	415954.2	NC	000913:1509286-1509462		VKQSE	FR		WLESQ	GVDV		ANGSN	HLKLR	FHGRR		VMPRHP		CD	41		
T10189	WP	000813794.1	NZ	CP025520:366728-366904		VKQSE	FR		WLESQ	GVDV		ANGSN	HLKLR	FHGRR		VMPRHP		CD	41		
T1325	WP	004079902.1	NC	000853:c1332998-1332765	MSYLP	IVDP	KTMEK		VLLKL	GFQR	VRQK	GSHV	FYRHS	NGKYT		TIPFHA		KE	49		
T875	15965792	NC	003047:2199475-2199663		VKSGD	IIA		ALQK	KDGYE	VATK	GSHV	QFKHP	KKHGR		VTV	PHP		KR	43		
T3593	YP	001085049.1	NC	009085:c2346189-2346031		MIEAD	GWYEV	RVS	GSHH	HF	KHP	TK	GL		VTI	PHP		KK	35		
T6107	WP	009893919.1	NC	006351:535977-536156		MNSSK	LIR		MLEED	GWRL	VRVT	GSHH	HF	KHP	KPG	L	VTI	PHP		KK	43
consensus						*	*	*				!	*	*	*	*	*	*	*	*	*

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T10122 AZV00182.1 MK291441:c9463-9056	IAHVALILDRGSPKRVNISLDPLGLDADAEARRGMT	RSFAFLSSAARAEIHAAH	135
T1039 NP 415954.2 NC 000913:1509286-1509462	ETKEPLRKAILKQL	GLS	58
T10189 WP 000813794.1 NZ CP025520:366728-366904	ETKEPLRKAILKQL	GLS	58
T1325 WP 004079902.1 NC 000853:c1332998-1332765	TCPSHLIRKIREA	CISVEEFKKVLENL	77
T875 15965792 NC 003047:2199475-2199663	DLPIGTLRSIEKQS	GLKLR	62
T3593 YP 001085049.1 NC 009085:c2346189-2346031	DLPNCTVKSIIKQA	GLN	52
T6107 WP 009893919.1 NC 006351:535977-536156	DLPIGTVKSIQKSA	GL	59
consensus	*** ** * ! * * * *	! * *	

- non conserved
- similar
- $\geq 50\%$ conserved