

logo

AT1049 NP 418645.2 NC 000913:4448447-4448698	.MRITIKRWGNSAGMVIPNIYMKELNLQPGQSV	EAQVSN	NQLITPTIS..RRYSIDELLA	AQCDMNAAEISEQDVW	72
AT1076 WP 000557619.1 NC 002134:91354-91611	MHTTRLKRVGGSVMLTVPPALLNALSIGTDNEVGMVIDN	GR	LIVEPYRR.PQYSLAELLA	AQCDPNAEISAEREW	74
AT10069 WP 011977835.1 NC 009653:1433-1663	.MTTATGKWNGAALRIPQPFMKQLGLSIGDVVNFRVTERDL	VITKAG....PTLEELL	QCTAE..NRHHEYLC		68
AT10130 WP 000581937.1 NZ HG738867:3872442-3872690	MIHSSVKRWGNSPA	VRIPATLMQALNLNIDDEVKIDLV	DCKLIIEPVRKEPVFTLAEV	VNDITPE..NLHENIDW	73
AT1044 NP 417263.1 NC 000913:c2911339-2911091	MIHSSVKRWGNSPA	VRIPATLMQALNLNIDDEVKIDLV	DCKLIIEPVRKEPVFTLAEV	VNDITPE..NLHENIDW	73
AT137 WP 010887061.1 NC 001263:417705-417947	.MTSQIQKWGNSLALRIPKALAQVGLTQSS	VEVLLQDGGQVIRPV	PAR.QYDLAALLAEMTPE..NLHGETDW		71
consensus	* * * * * ! * * * * * ! * * * * * ! * ! * * * * * * * * * * * ! * * ! * * ! * * * * * * * * * * * * *				

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AT1049 NP 418645.2 NC 000913:4448447-4448698	GKSTPACDEIW	83
AT1076 WP 000557619.1 NC 002134:91354-91611	LDAPATCQEEI	85
AT10069 WP 011977835.1 NC 009653:1433-1663	D...SQCKENL	76
AT10130 WP 000581937.1 NZ HG738867:3872442-3872690	GE..PKDKEVW	82
AT1044 NP 417263.1 NC 000913:c2911339-2911091	GE..PKDKEVW	82
AT137 WP 010887061.1 NC 001263:417705-417947	GA..LEGREEW	80
consensus	* * * * ! * *	

- X non conserved
- X similar
- X ≥ 50% conserved