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T111	NP	216619.1	NC	000962:c2364520-2364086	.MKIVDANVLIYAVNTTSEHHKPSLRWIDGALSGADRVGFAWVPLLAFVRLATKVGLFPRPLPREAAITQVAD..WL...	74
T119	NP	217118.1	NC	000962:2930344-2930784	.MLLCDTNITWLALALSGHVHHRASRAWLDTINAPG.VIHFCRATQQSLRLLTNRTVLGAYGSPPLTNR....EAWAAYA	74
T80	15607757	NC	000962:711006-711407		VTVLLDANVLIALVVAEHVHHDAAADWLMA.SDTG.F.ATCPMTQGS�VRFLVR....SGQSAAAARDVVSAVQCTSRH	72
T129	NP	217837.1	NC	000962:c3708070-3707642	MRALLDVNVLLALLDRDHVDHERARAWITGQIERG.W.ASCAITQNGFVRVISQPR.YPSPISVAHAIIDLARATHTRYH	77
T112	NP	217010.1	NC	000962:2808310-2808735	.VALLDVNALVALAWDSHTHHARIREWFTANATLG.W.ATCPLTEAGFVRVSTNPKVLPSAIGIADARRVLVALRAVGGH	77
T114	NP	217046.1	NC	000962:c2854686-2854267	VTALLDVNVLIALGWPNHVHHAAAQRWFTQFSSNG.W.ATTPITEAGYVRISSNRSVMQVSTTPAIAIAQLAAMTSLAGH	78
consensus					***! !***** ***! !* * * * * * * * * * * * * * *	

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T111	NP	216619.1	NC	000962:c2364520-2364086	...AAPSAVLVNPTVRHADI.LARMLTYVGTGANLVNDAHLAALAVEHRASTIVSYDSDFGRFEGVRWDQPPALL...	144
T119	NP	217118.1	NC	000962:2930344-2930784	AFLLDDRIVLAGAEPDGLEAQWRAFAVRQSPAPKVWMDAYLAFAFALTGGFELVTTDTAFTQYGGIELRLLAK.....	146
T80	15607757	NC	000962:711006-711407		EFWPDALSFAGV.....EVAGVVGHRQVTDAYLAQLARSHDQQLATLDSGLAHLHGDVAVLIPT..TT.	133
T129	NP	217837.1	NC	000962:c3708070-3707642	EFWSCTVSIILDSKV.....IDRSRLHSPKQVTDAYLLALAVAHDGRFVTFDQSIALTAVPGATKQ..HLATL	142
T112	NP	217010.1	NC	000962:2808310-2808735	RFLADDVSLVDD.....DVPLIVGYRQVTDHLLTLARRRGVRLVTFDAGVFTLAQQRPKTPVELLTIL	141
T114	NP	217046.1	NC	000962:c2854686-2854267	TFWPDVPLIVGSA.....GDRDAVSNHRVTDCHLIALAARYGRLVTFDAALADSASAG...LVEVL...	139
consensus					* *	

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