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*:*****:*****:* .:.* ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** ** **
gi | 4502987 | ref | NP_001855.1 | MQALRVSQALIRSFSSSTARNRVREKQKLFQEDNDIPLYLKGGIVDNILYRVTMTLCLGGTVVSYSLGWASFPNRN- 79
gi | 73948429 | ref | XP_855440.1 | MRALRVSQALVRSFSSSTARNRVQNRVAEKQKLFQADNDLPVHLKGGGTDNILYRVTMGLCLGGTVVSYFLGWASFPHKK 80
gi | 28603808 | ref | NP_788847.1 | MRALRVSQALVRSFSSSTARNRVFNVAEKQKLFQEDNGLPVHLKGGATDNILYRVTMTLCLGGTLYSLYCLGWASFPHKK 80
gi | 6753504 | ref | NP_034074.1 | MRALRVSQALVRSFSSSTRSHLENRVAEKQKLFQADNDLPVHLKGGGMDNVLYRLTMTLTLGGTAYCLYCLGWASFPHKK 80
1.....10.....20.....30.....40.....50.....60.....70.....80

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