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gi|9624979|ref|NP_062507.1|MSQKQEEENPAEETGEEKQDTQEKEGILPEKAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGADKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|11177874|ref|NP_068614.1|MSQKQEEENPAEETGEEKQDTQEKEGILPEKAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGADKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|27806639|ref|NP_776475.1|MSQKQEEENPAEETGEEKQDTQEKEGILPEKAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGPDKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|114559309|ref|XP_001169491.1|MSQKQEEENPAEETGEEKQDTQEKEGILPERAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGPDKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|4758272|ref|NP_004427.1|MSQKQEEENPAEETGEEKQDTQEKEGILPERAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGPDKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|73981333|ref|XP_860994.1|MSQKQEEENPAEETGEEKQDTQEKEGILPERAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-SAGPDKNLVTGDHIPTPODLPQRKSSSLVTSKLAGGQVE 122
gi|56118972|ref|NP_001007966.1|MAAPLGTGARAEDSGQEKQDSQEKETVIPERAEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLP-TAGPDKNLVTGDHIFKPPDLPQRKSSSLVASKLAG---- 122
gi|292622386|ref|XP_002664958.1|MSSDNDTETHPEQVDETESQDTN-SNPVRETEAAKLRKAKYPSLGQKPGGSDFLMKRLQKQKYFDSGDYNMAKAKMKNKQLPAAAGPDKNIVTGDHIPTPODLPQRKSSSLVTSKLAG---- 122
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120..

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