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gi | 83367075 | ref | NP_783316.2 | MDPNCS CATGG SCTCAGS CKCKECKCTS CKKS CCS CCPVGC AKCAQGC VCKGASEKCS CCA 61
gi | 114664338 | ref | XP_001144626.1 | MDPNCS CATGG SCTCAGS CKCKECKCTS CKKS CCS CCPVGC AKCAQGC VCKGASEKCS CCA 61
gi | 55741622 | ref | NP_001003173.1 | MDPDCS CSTGG SCTCAGS CKCKECKCTS CKKS CCS CCPVGC AKCAQGC ICKGASDKCS CCA 61
gi | 7305285 | ref | NP_038630.1 | MDPNCS CSTGG SCTCTSS CACKNCKCTS CKKS CCS CCPVGC SKCAQGC VCKGAADKCTCCA 61
gi | 20301948 | ref | NP_620181.1 | MDPNCS CSTGG SCTCSS CGCKNCKCTS CKKS CCS CCPVGC SKCAQGC VCKGASDKCTCCA 61
gi | 94966789 | ref | NP_001035582.1 | MDPNCS CPTGG SCS CAGS CTCKACRCPS CKKS CCS CCPVGC AKCAQGC VCKGASDKCS CCA 61
gi | 118151342 | ref | NP_001071602.1 | MDPNCS CPTSG SCS CAGS CTCKACRCPS CKKS CCS CCPVGC AKCAQGC ICKGASDKCRCHA 61

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1.....10.....20.....30.....40.....50.....60.

