

gi|51479166|ref|NP_001003800.1| 150
gi|114625574|ref|XP_001147166.1| 150
gi|85702355|ref|NP_001034268.1| 150
gi|76008363|ref|NP_001028846.1| 150
gi|73947197|ref|XP_541330.2| 150
gi|118096856|ref|XP_414332.2| 150
gi|68442045|ref|XP_685529.1| 150



gi|51479166|ref|NP_001003800.1| 298
gi|114625574|ref|XP_001147166.1| 298
gi|85702355|ref|NP_001034268.1| 300
gi|76008363|ref|NP_001028846.1| 300
gi|73947197|ref|XP_541330.2| 291
gi|118096856|ref|XP_414332.2| 296
gi|68442045|ref|XP_685529.1| 294



gi|51479166|ref|NP_001003800.1| 445
gi|114625574|ref|XP_001147166.1| 445
gi|85702355|ref|NP_001034268.1| 447
gi|76008363|ref|NP_001028846.1| 447
gi|73947197|ref|XP_541330.2| 438
gi|118096856|ref|XP_414332.2| 444
gi|68442045|ref|XP_685529.1| 442



gi|51479166|ref|NP_001003800.1| 595
gi|114625574|ref|XP_001147166.1| 595
gi|85702355|ref|NP_001034268.1| 591
gi|76008363|ref|NP_001028846.1| 591
gi|73947197|ref|XP_541330.2| 581
gi|118096856|ref|XP_414332.2| 586
gi|68442045|ref|XP_685529.1| 575



gi|51479166|ref|NP_001003800.1| 745
gi|114625574|ref|XP_001147166.1| 745
gi|85702355|ref|NP_001034268.1| 741
gi|76008363|ref|NP_001028846.1| 741
gi|73947197|ref|XP_541330.2| 726
gi|118096856|ref|XP_414332.2| 736
gi|68442045|ref|XP_685529.1| 719



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*****:***:***** **:*:*:*
gi | 51479166 | ref | NP_001003800.1 | LRAMFATRCDEYITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELLELDHEQTRRGRAKAAPKTKPATPSVSHTCACASDRAEGTGLANQVFCSEKHSIYCD 855
gi | 114625574 | ref | XP_001147166.1 | LRAMFATRCDEYITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELLELDHEQTRRGRAKAAPKTKPVTPSVSHTCACASDRAEGTGLANQVFCSEKHSIYCD 855
gi | 85702355 | ref | NP_001034268.1 | LRAMFATRCDEYITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELLELDHEQTRRGRSKAASKAKPASPSVSHTCACASERAEGAGLANQVFCSEKHSIYCD 851
gi | 76008363 | ref | NP_001028846.1 | LRAMFATRCDEYITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELLELDHEQTRRGRNKATSKAKPATPSVSHTCACASERAEGTGLANQVFCSEKHSIYCD 851
gi | 73947197 | ref | XP_541330.2 | LRAMFATRCDEYITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELLELDHEQTRRGRAKATSKAKPGTSPVSHTCACASDRAEGAGLITVLCGEKYNIIYCD 836
gi | 118096856 | ref | XP_414332.2 | LRAMFATRCDEYVITQLDEMORQLAAAEDEKKTLSNLLRMAIQKLALTRLEHLELDHEQSKRVRTKSASKAKSSNPITL-----GRGRPLSFSSPH----- 815
gi | 68442045 | ref | XP_685529.1 | LRAMFATRCDEYVITQLDDMORQLAAAEDEKKTLSNLLRMAIQKLALTRLELDLEFDNEQSRRC-SGAGPGGRGKAPS-----GRGRPLSFSSPH----- 809
.....760.....770.....780.....790.....800.....810.....820.....830.....840.....850.....860

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