

gi | 225690502 | ref | NP_001139500.1 | --MSPGGKFD... 150
gi | 73962639 | ref | XP_547737.2 | --MSPGGKFD... 150
gi | 194670481 | ref | XP_001249446.2 | --MSPGGKFD... 150
gi | 31982212 | ref | NP_796023.2 | --MSPGGKFD... 150
gi | 51172604 | ref | NP_001003711.1 | MHVPLGRKFD... 150



gi | 225690502 | ref | NP_001139500.1 | YHQAALLRSPRRTSLD... 300
gi | 73962639 | ref | XP_547737.2 | YHQAALLRSPRRTSLD... 300
gi | 194670481 | ref | XP_001249446.2 | YHQAALLRSPRRTSLD... 300
gi | 31982212 | ref | NP_796023.2 | YHQAALLRSPRRTSLD... 300
gi | 51172604 | ref | NP_001003711.1 | YHQAALLRSPRRTSLD... 300



gi | 225690502 | ref | NP_001139500.1 | RSNGLRYEGEWLGNRRHGYGRTRPDGSRREEGKYKRNRLVHGGRVRSLLPLALRRGKVKVDRAVEGARRAVSAARORQ... 450
gi | 73962639 | ref | XP_547737.2 | RSNGLRYEGEWLGNRRHGYGRTRPDGSRREEGKYKRNRLVHGGRVRSLLPLALRRGKVKVDRAVEGARRAVSAARORQ... 450
gi | 194670481 | ref | XP_001249446.2 | RSNGLRYEGEWLGNRRHGYGRTRPDGSRREEGKYKRNRLVHGGRVRSLLPLALRRGKVKVDRAVEGARRAVSAARORQ... 450
gi | 31982212 | ref | NP_796023.2 | RSNGLRYEGEWLGNRRHGYGRTRPDGSRREEGKYKRNRLVHGGRVRSLLPLALRRGKVKVDRAVEGARRAVSAARORQ... 450
gi | 51172604 | ref | NP_001003711.1 | RSNGLRYEGEWLGNRRHGYGRTRPDGSRREEGKYKRNRLVHGGRVRSLLPLALRRGKVKVDRAVEGARRAVSAARORQ... 450



gi | 225690502 | ref | NP_001139500.1 | NGLTPSEGSPPELSPASSRQ... 600
gi | 73962639 | ref | XP_547737.2 | NGLTPSEGSPPELSPASSRQ... 600
gi | 194670481 | ref | XP_001249446.2 | NGLTPSEGSPPELSPASSRQ... 600
gi | 31982212 | ref | NP_796023.2 | NGLTPSEGSPPELSPASSRQ... 600
gi | 51172604 | ref | NP_001003711.1 | NGLTPSEGSPPELSPASSRQ... 600



gi | 225690502 | ref | NP_001139500.1 | AOPGAANPLVVGAVALLDL... 630
gi | 73962639 | ref | XP_547737.2 | AOPGAANPLVVGAVALLDL... 630
gi | 194670481 | ref | XP_001249446.2 | AOPGAANPLVVGAVALLDL... 630
gi | 31982212 | ref | NP_796023.2 | AOPGAANPLVVGAVALLDL... 630
gi | 51172604 | ref | NP_001003711.1 | AOPGAANPLVVGAVALLDL... 630

