

gi	ref	seq	pos
gi 70887541	ref NP_034958.2	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 16758512	ref NP_446150.1	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 19923343	ref NP_006070.2	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 114609541	ref XP_518770.2	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 115495013	ref NP_001069287.1	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 73945615	ref XP_541127.2	MADHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 45598375	ref NP_996726.1	MAEHMMAMNHGRFPDGTNGLHHHPAHRMGMGQFPSPHHHQQQQPO-HAFNALMGEHIIHYGAGNMNATS	1
gi 54400592	ref NP_001006045.1	MVDRMAMNHGRFPDAVNG-HQHSARFMGMGQFSNALPQQG-----HYNVIMGDHMHYAGGNIINANNIRHSMATGNIN	1



gi	ref	seq	pos
gi 70887541	ref NP_034958.2	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 16758512	ref NP_446150.1	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 19923343	ref NP_006070.2	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 114609541	ref XP_518770.2	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 115495013	ref NP_001069287.1	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 73945615	ref XP_541127.2	NGTNQHFRDCNPKHSGGSSIPGGAGGSGTTPGGSGGTSGGAGGSSAGGSGGG---ETMPASVAHVPAAMLPPNVIDDFIDEEVLM	210
gi 45598375	ref NP_996726.1	SGGGQHFRDCNPKHGGG-----GGG-----LPPAVPHVPAAMLPPNVIDDFIDEEVLM	210
gi 54400592	ref NP_001006045.1	NGTGGQFRDSIAKQNTSG-----VLPGHMHPAAMLPPNVIDDFIDEEVLM	210

