

gi	6755674 ref NP_035620.1	LPMVPEVAQAVGVSQGHHTKDFTR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	16758178 ref NP_445888.1	LPMVPEVAQAVGVSQGHHTKDFAR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	82659087 ref NP_059347.2	LPMVPEVAQAVGVSQGHHTKDFTR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	114682565 ref XP_514710.2	LPMVPEVAQAVGVSQGHHTKDFTR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	73992231 ref XP_867111.1	LPMVPEVAQAVGVSQGHHTKDFTR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	119906050 ref XP_591710.3	LPMVPEVAQAVGVSQGHHTKDFAR-VAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	61098218 ref NP_001012849.1	LPMVPEVAQAVGASQGHHTKEFNR-AAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	45387553 ref NP_991124.1	LPMVPEVAQAVGANQGPFAKDYGRTAAPNPAKATVTAMIARELLYGGTSPTAETILKSNISSGHVPHGPRTRPSEQLYLLSRAQGFQVEYKDFPKNNKNECVSLINCSSQPPLVSHGIGKDVESCHDMAALNILKLLSELDQOSTEMPRT	750
gi	17533821 ref NP_494501.1	-----QS-----	750
gi	115534938 ref NP_508195.2	-----STELTDINEKQKVMEKVAHCLSEFEIITDLN-----	750
gi	17568157 ref NP_508196.1	PPGDQDYFNLSIVQSLKKDAIVEGKIRRLKRSKENRRALTAELIVEL---ERAQSYLQTKNTIQSSQSSAHHHLEQLSDFKFSLOYTSFPQVGDQHFIVSIGLEAPLVGHGTCCSTTEADENAALDAIKLELSASKI-----	750
	610.....620.....630.....640.....650.....660.....670.....680.....690.....700.....710.....720.....730.....740.....750	



gi	6755674 ref NP_035620.1	GNGPVSACGRC-----	801
gi	16758178 ref NP_445888.1	GNGPVSACGRC-----	801
gi	82659087 ref NP_059347.2	GNGPMSVCGRC-----	801
gi	114682565 ref XP_514710.2	GNGPMSVCGRC-----	801
gi	73992231 ref XP_867111.1	GNGPMSVCGRC-----	801
gi	119906050 ref XP_591710.3	GNGPMSVCGRC-----	801
gi	61098218 ref NP_001012849.1	GNGPMSVEVKIPRAVTFPLRRTRGCVKQEMESDPLLKPAANSNITLGGTLDSTA	801
gi	45387553 ref NP_991124.1	ANGPMSGCTKQDLEG-----DALLKPAANSSTMGKTLDDGTA	801
gi	17533821 ref NP_494501.1	-----	801
gi	115534938 ref NP_508195.2	-----	801
gi	17568157 ref NP_508196.1	-----	801
	760.....770.....780.....790.....800.	

