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gi |45551086|ref|NP_725084.2| CGHCKKLSPIVEELAEKLQDE-DVAIVKMDAANDVPPPEFNVRGFPTLFWLPKDAKNKPVSYNGGREVDDFLKYLAKKATTELKGFDR---SGKPKK-TEL 489
gi |158285742|ref|XP_001687942.1| CGHCKKLTPTLEELGTGLKDE-AVSIKMDAANDVPPQFEVRFPTLFWLPKDAKSSPARYEGGREVDDFKYIAKHAISELKGFDL---AGGAKK-TEL 488
gi |17507915|ref|NP_491995.1| CGHCKSLAPKYVEELAEKLNKE-DVVIKMDAANDVPPMFEVRFPTLFWLPKNAKSNPIYNGGREVKDFVYFISKHSIDGLKGFDR---DGKKKKTEL 488
gi |57526015|ref|NP_001003517.1| CGHCKKLEPKYTALGEMLYSDPNIVIAKMDAANDVPPAGYDVGFPPIYFAAAGRKSEPKRYEGAREVKDFVNFVKREAIKPLILNGV-----KDEL 493
gi |125840013|ref|XP_001331697.1| CGHCKNLEPKYKELGKLSGNPNIVIAKMDAANDVPPNVDVGFPTIYFVPSGQKQDPRRYEGGREVNDFIYLYKKEATNPLILDDS-----RDEL 488
gi |112293264|ref|NP_031978.2| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLTPKKYEGGRELNDLISYLQREATNPPVIOEE-KPKK-KKKAQEDL 505
gi |8393322|ref|NP_059015.1| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLTPKKYEGGRELNDLISYLQREATNPPVIOEE-KPKK-KKKAQEDL 505
gi |21361657|ref|NP_005304.3| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLNPKKYEGGRELSDFISYLQREATNPPVIOEE-KPKK-KKKAQEDL 505
gi |114656683|ref|XP_001159572.1| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLNPKKYEGGRELSDFISYLQREATNPPVIOEE-KPKK-KKKAQEDL 505
gi |148230374|ref|NP_776758.2| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLNPKKYEGGRELSDFISYLQREATNPPVIOEE-KPKK-KKKAQEDL 505
gi |74000349|ref|XP_535453.2| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFSPANKKLNPKKYEGGRELSDFISYLQREATNPPVIOEE-KAKK-KKKAQEDL 687
gi |45383890|ref|NP_989441.1| CGHCKNLEPKYKELGKLSKDPNIVIAKMDAANDVPPSPYEVKGFPTIYFAPAGKQSPKKYEGGREVSDFISYLQREATSTPVLOEDKAKKSKKKAQEDL 505
gi |115446679|ref|NP_001047119.1| CGHCKLAPILLEVAVSLKDEEDVVIKMDAANDVPPSDFAVEGYPSMYFYSSGNLLEFYDGRTAEEIIDFIKKNKGSRPGEATTES-----VKDEL 545
.....610.....620.....630.....640.....650.....660.....670.....680.....690.....700..

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