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gi|6681233|ref|NP_031911.1| KNLREKSGIKVFRFIAPLYYINKECFKSALYKKA-LNPVLVKAAWKKAAR-RKLKEEMVTFRGGDPDEVSMQLSHDPLEVHTIVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLARGEYCK--KEEENLLFYSLSEA 750
gi|16923978|ref|NP_476468.1| KNLREKSGIKVFRFIAPLYYINKECFKSALYKKT-LNPVLVKAAWKKAAR-RKLKEETVTFHGGDPDEVSMQLSHDPLELHTVVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLARGEYCK--KEEENLLFYSLSEA 750
gi|100913030|ref|NP_000103.2| KNLQIKPGIKIFRFVAPLYYINKECFKSALYKQT-VNPILIKVAWKKAAR-RRIKEKVVTLGGIDDEMSVQLSHDPLELHTIVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLINGEYCK--KEEENLLFYSVYEA 750
gi|114602780|ref|XP_001164642.1| KNLQIKPGIKIFRFVAPLYYINKECFKSALYKQT-VNPILIKVAWKKAAR-RRIKEKVVTLGGIDDEMSVQLSHDPLELHTIVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLINGEYCK--KEEENLLFYSVYEA 750
gi|73954204|ref|XP_855107.1| KNLQIKPGIKIFRFVAPLYYINKECFKSALYKKT-LNPVLVKAQKKAAR-RKITKETVILSGVQDEVSVQLSHDPLELQIVVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLARGEYCK--KEEENLLFYSVYEA 750
gi|94966855|ref|NP_001035615.1| KNLQAKSGIKIFRFVAPLYYINKECFKSALYKKT-LNPVLVKAQKKAAR-RKIKRETVPSSGIDDEVSVQLSHDPLEFHTIVIDCSAIQFLDTAGIHTLKEVRRDYEAVGIQVLLAOCNPSVRDSLARGEYCK--KDEENLLFYSIYEA 750
gi|118097311|ref|XP_425183.2| KNLQIKPGVVVLRFEAPLYYINKECFKSALYKQTGVNPAALVKAQKKAAR-RMLREKEAGSGGNGTSSISMELVSEPLGFHTIVIDCCAVQFLDTAGIHTLKEVCKDYNEIDVQVLLAOCNPSVRSRLMRGEFFK--EGEDHLLFHSVHQA 750
gi|68373377|ref|XP_685114.1| NGLQIQPDVAVFRNEAPIYYANQSLFKKSLYRNVLGDLPLKEKAKQKLEKQRKQKHSREDQKQEMDASTNVNLLQHSFHTLVIDCSVLFLLDTAGVNALKEVFKDYKELGVNLLAOCNPSVIDSLRRGDYDFNKGTKDIQFHITGEA 750
gi|15233255|ref|NP_188220.1| REASRIPGFLILATESPIYFANSTYLDRLRWA-----REEENRIKE-----NNGTTLKCIILDMTAVSAIDTSGLEAVFELRRRLEKQSLQLVLVNPVGTVMKELHKSIIIE--ALGLSGLYLVVGEA 750
gi|115466358|ref|NP_001056778.1| REAMRVPSTLVVGVESAIYFANSMYLGRIIMRFL-----REEDERAAR-----CNQCQVRCIILDMSAVAIDTSGLDALAEKLVLEKRNIELVLANPVGEVTERLYNSVVGK--TFGSDRVFFSVAEA 750
.....610.....620.....630.....640.....650.....660.....670.....680.....690.....700.....710.....720.....730.....740.....750

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gi|6681233|ref|NP_031911.1| VAFAE+DS+ON+Q+KG+VC+VV+N+GL+SL+SG+D 774
gi|16923978|ref|NP_476468.1| VAFAE+ES+Q+EK+KG+VC+VV+N+GL+SL+SG+D 774
gi|100913030|ref|NP_000103.2| MAFAE+VS+K+N+Q+KG+VC+VP+N+GL+SL+SS+D 774
gi|114602780|ref|XP_001164642.1| MAFAE+VS+K+N+Q+KG+VC+VP+N+GL+SL+SS+D 774
gi|73954204|ref|XP_855107.1| MAFAE+ES+ON+Q+KG+VC+IP+N+GL+SL+SS+D 774
gi|94966855|ref|NP_001035615.1| M+FAE+DS+ON+Q+KE+R+H+IP+NG+NE+FS+D 774
gi|118097311|ref|XP_425183.2| VDFALGA+EH+SRI+CASKN----- 774
gi|68373377|ref|XP_685114.1| ILYG+SL+KS+NG+NC+DT+SV----- 774
gi|15233255|ref|NP_188220.1| VADLSS+WKANGQP----- 774
gi|115466358|ref|NP_001056778.1| VAAAP+---HK+TQP----- 774
.....760.....770....

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