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gi | 24662483 |ref | NP_648435.1 | LSECNIPGEC--IVEQDI GRLKSI AAKMLSD LGMH- A IISDDV LHEI -----CRYGGAELH AVS AF IGGCAAQ EVIK IITK QYKPIDNT F IYN AIT TE E VTLK L 524
gi | 58387417 |ref | XP_315544.2 | QADYGC L PGEH --DPESD I TGMKSI AAKMLSD WGIG- PPLSDDLAYEI -----CRYGGAEIHS I SA YLGGCIAHELIKLVTRQYRPF DNT F IYDGS T SQ EET YKL 523
gi | 57524906 |ref | NP_001006129.1 | VKOHGRYPGVYNYQVEDDI GKLSCLT SFLOEHGLS- VLVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 535
gi | 118123592 |ref | XP_001236466.1 | VKOHGRYPGVYNYQVEDDI GKLSCLT SFLOEHGLS- VLVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 517
gi | 21450341 |ref | NP_659180.1 | HKOHGRYPGVSNYQVEDI GKLSCLT GFLOEYGLS- VMVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 534
gi | 14091760 |ref | NP_114461.1 | HKOHGRYPGVSNYQVEDI GKLSCLT GFLOEYGLS- VMVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 534
gi | 4502169 |ref | NP_003896.1 | HKOQGRYPGVSNYQVEDI GKLSCLT GFLOEYGLS- VMVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 534
gi | 114663003 |ref | XP_001159481.1 | HKOHGRYPGVSNYQVEDI GKLSCLT GFLOEYGLS- VMVKDDYVHELLIKIHVS VVSFS CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 546
gi | 73957330 |ref | XP_852003.1 | HKOHGRYPGVSNYQVEDI GKLSCLT GFLOEYGLS- VMVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 534
gi | 119910184 |ref | XP_871117.2 | HKOHGRYPGVSNYQVEDI GKLSCLT SFLOEYGLS- VMVKDDYVHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 534
gi | 41053883 |ref | NP_956793.1 | YQDHSRYPGVYNYQVEDI INK LKLVN SLLQ EYSLN- VMVKDDYIHEF -----CRYGAAEPHT IAAFLGGAAAQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 533
gi | 15225241 |ref | NP_180800.1 | AGT YKKFPGQFDGSD EDASQLKTIALSLLS E MCGDGV ELQ EELYNEM -----CRFGAAEIHVVAALIGGITSQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 523
gi | 15220442 |ref | NP_172010.1 | AANYNKFPQFDGMD EDSRLK TIALSLLTDLGCNGS VLPDDL IHEM -----CRFGASEIHVVSAFVGGIASQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 540
gi | 115456233 |ref | NP_001051717.1 | AANYNRCPGIFESIEDV PRLKTA AVS VMS EMMNGAPL SEDL IHEM -----CRFGAAEIHVVAALIGGIVASQ EVIK IITK QFVIFNNT Y IYSGMSQTSATFQL 531
.....460.....470.....480.....490.....500.....510.....520.....530.....540.....550.....560..

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