

gi | 4506401 | ref | NP_002871.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 114585472 | ref | XP_001155232.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 18497290 | ref | NP_084056.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 6981458 | ref | NP_036771.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 73984570 | ref | XP_859234.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 156523122 | ref | NP_001095975.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 45384314 | ref | NP_990638.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150
gi | 40538760 | ref | NP_571514.1 | MEHIQGAWKITISNGFGPKDAVDFDSSCIPTIVQVQFGYORRASDDGKLTDPSSKTSNTIRVFLPNKQRTVVNVRNGMSLHDCLMKALKVVRGLOPECCAVFRLLEHKGKARLDWNTDAASLIGEELOVDFLDHVPLTTHNFARKTFLKLA 150



gi | 4506401 | ref | NP_002871.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 114585472 | ref | XP_001155232.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 18497290 | ref | NP_084056.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 6981458 | ref | NP_036771.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 73984570 | ref | XP_859234.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 156523122 | ref | NP_001095975.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 45384314 | ref | NP_990638.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300
gi | 40538760 | ref | NP_571514.1 | FCDICQKFLNNGFRQTCGYKFFHEHCSTKVPIMCVDWNSNIRQLLFPNSTIGDSGVPALPPLTMRMRRESVSRMPVSSQHRYSHPHAFIFNTSSPSSSEGLSQRORSTSTPNVHMVSTLTPVDSRMIE-----DA 300



gi | 4506401 | ref | NP_002871.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 114585472 | ref | XP_001155232.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 18497290 | ref | NP_084056.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 6981458 | ref | NP_036771.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 73984570 | ref | XP_859234.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 156523122 | ref | NP_001095975.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 45384314 | ref | NP_990638.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450
gi | 40538760 | ref | NP_571514.1 | IRSHSESASPALSSSPNNLSPITGWSQPKTPVPAQRERAPVSGTQEKNKIRPRGQDSSYYWEIEASEVMLSTRIGSGSPGTVYKKGWHDVAVKILKVVDPPTPEQFAFRNEVAVLRKTRHVNILLFPMGYMTKDNLAIVTQWCEGSSLY 450



gi | 4506401 | ref | NP_002871.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 114585472 | ref | XP_001155232.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 18497290 | ref | NP_084056.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 6981458 | ref | NP_036771.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 73984570 | ref | XP_859234.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 156523122 | ref | NP_001095975.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 45384314 | ref | NP_990638.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600
gi | 40538760 | ref | NP_571514.1 | KHLHVQETKFMFLIDDIARQTAQGM DYLHAKNIIHRDMKSNIFLHEGLTVKIGDFGLATVKSRSWGSQOQVEOPTGSLVWMAPEVIRMODNPPSFDQSDVYSYGVIVLYELMTGELPYSHINNRDQIIFMVGRGYASFDLSKLYKNCPKA 600



gi | 4506401 | ref | NP_002871.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 114585472 | ref | XP_001155232.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 18497290 | ref | NP_084056.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 6981458 | ref | NP_036771.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 73984570 | ref | XP_859234.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 156523122 | ref | NP_001095975.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 45384314 | ref | NP_990638.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668
gi | 40538760 | ref | NP_571514.1 | MKRLVADVCKKVKERPLFPQILSSIELLOHSLPKINRSASEPSLHRAAHTEDINACTLTTSPRLPVF 668

