

gi | 195976807 | ref | NP_002890.2 | -----MDPPAGFVVRAGNPAVAAPQSPLSPGAFHRAAHHPRTGSRCPGSLQPSRPLVANWLSLPEMPVDFGTGWKMLVNE--FEEYLRALDNNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
gi | 114589486 | ref | XP_001158395.1 | -----MDPPAGFVVRAGNPAVAAPQSPPSPGAFHRAAHHPRTDRCRCPGSLQPSRPLVANWLLPEMPVDFGTGWKMLVNE--FEEYLRALDNNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
gi | 70778780 | ref | NP_001020514.1 | -----MPVDFGTGWKMLVNE--FEEYLRALDNNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
gi | 6981464 | ref | NP_036865.1 | -----MPVDFNGYWKMLSNEN--FEEYLRALDNNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
gi | 6755300 | ref | NP_035384.1 | -----MPVDFNGYWKMLSNEN--FEEYLRALDNNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
gi | 50752060 | ref | XP_422635.1 | -----MPADFNQYWKMLSNEN--FEEYLKALDNNVAVRRIANLLKPKDKEILQGDHMIIRTLSTFRNYIMEFVVGKE 150
gi | 168823474 | ref | NP_001108371.1 | -----MSPVDMNGYWKMLSNEN--FDEYMEALDNNVAVRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMEFVVGKE 150
gi | 73990638 | ref | XP_534285.2 | -----MLSPATLDHPFPEASLSVASLLQLVSLPLQSVLYQLGRANLYPLPLLSVPLNCSQSVQDTLCTSLCSLCPHLPYTCPPPGGTSRAVSSASFPVPGTCQVLNVALRRIANLLKPKDKEIVQGDHMIIRTLSTFRNYIMDFVVGKE 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150



*****:*****:*** *****:*****:***:***:*** ** *****:
gi | 195976807 | ref | NP_002890.2 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVQ-- 217
gi | 114589486 | ref | XP_001158395.1 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVQ-- 217
gi | 70778780 | ref | NP_001020514.1 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVN-- 217
gi | 6981464 | ref | NP_036865.1 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVH-- 217
gi | 6755300 | ref | NP_035384.1 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVV-- 217
gi | 50752060 | ref | XP_422635.1 | FEEDLAGVDDRKCMITVSWDGDKLLCVQKGEKEDRGWTQWIEGDEMHLIRVCGVCKQVFKKVQ-- 217
gi | 168823474 | ref | NP_001108371.1 | FEEDLVGVDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWVKGDELHVMRVCVGVCKQVFKRQVFD 217
gi | 73990638 | ref | XP_534285.2 | FEEDLTGIDDRKCMITVSWDGDKLCVQKGEKEGRGWTQWIEGDELHLEMRVCGVCKQVFKKVQ-- 217
.....160.....170.....180.....190.....200.....210.....

