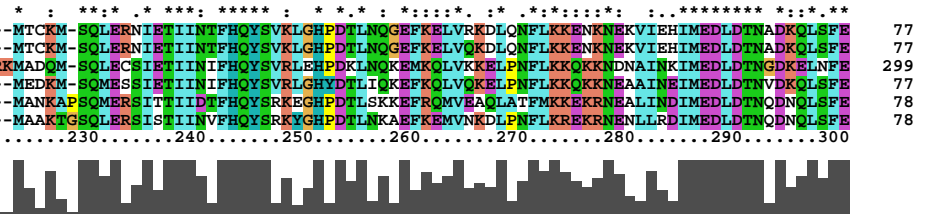


gi | 4506773 | ref | NP_002956.1 |
gi | 114559681 | ref | XP_001137562.1 |
gi | 73961682 | ref | XP_547585.2 | MAPNHAIAGGSELPGEDPFPYHPTSQPVALVGSEDRWHWGSDWMEPEISCLGQDLHSSVPAAPDQRPASLEHYGLLMQENHYOQGTHRPEGDVITPGLLAPLSPRLLSARTSAVKECLBOSQVGEKEGLLDGWHAGEMOSQOPLTDF 150
gi | 114052490 | ref | NP_001039793.1 |
gi | 6677837 | ref | NP_033140.1 |
gi | 16758364 | ref | NP_446039.1 |
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

gi | 4506773 | ref | NP_002956.1 |
gi | 114559681 | ref | XP_001137562.1 |
gi | 73961682 | ref | XP_547585.2 |
gi | 114052490 | ref | NP_001039793.1 |
gi | 6677837 | ref | NP_033140.1 |
gi | 16758364 | ref | NP_446039.1 |
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....290.....300



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gi | 114559681 | ref | XP_001137562.1 |
gi | 73961682 | ref | XP_547585.2 |
gi | 114052490 | ref | NP_001039793.1 |
gi | 6677837 | ref | NP_033140.1 |
gi | 16758364 | ref | NP_446039.1 |
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