

```

gi|7106315|ref|NP_032032.1|-----MAAGSITSLPALPEDGG 150
gi|9506599|ref|NP_062178.1|-----MAAGSITSLPALPEDGG 150
gi|153285461|ref|NP_001997.5|MVGVGGDVEDVTPRPGGCISSGRGARGCNGIPGAAAWAALPRRRPRRHPSVNPSSRAAGSPRIIRGRRTTEERPSSGRLGDRGRGRALPGGRLGGRGRGRAPERVGGGRGRGRTAAPRAAPAARGSRPGPAGTMAAGSITSLPALPEDGG 150
gi|27806627|ref|NP_776481.1|-----MAAGSITSLPALPEDGG 150
gi|45382175|ref|NP_990764.1|-----MAAGAAGSITSLPALPEDGG 150
gi|47086387|ref|NP_997988.1|-----MATGGITSLPAAP-DAE 150
1.....10.....20.....30.....40.....50.....60.....70.....80.....90.....100.....110.....120.....130.....140.....150

```



```

..**.*:*****:*****:***** * * ***:..**.*: ***: * .*:**:*:*****:*****.***** **:*:*****.
gi|7106315|ref|NP_032032.1|-AAPPGHFKDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHVKLQLOAERGVVSIKGVCANRYLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSSWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
gi|9506599|ref|NP_062178.1|-GAPPGHFKDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHVKLQLOAERGVVSIKGVCANRYLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSSWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
gi|153285461|ref|NP_001997.5|SGAPPGHFKDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHIKLQLOAERGVVSIKGVCANRYLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSSWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
gi|27806627|ref|NP_776481.1|SGAPPGHFKDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHIKLQLOAERGVVSIKGVCANRYLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSSWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
gi|45382175|ref|NP_990764.1|GGAPPGHFKDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHIKLQLOAERGVVSIKGVSANRFLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSDWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
gi|47086387|ref|NP_997988.1|NSAPPAGSFRDPKRLYCKNGGFLLRIHPDGRVDGVREKSDPHIKLQLOAERGVVSIKGVSANRFLAMKEDGRLLSKCVTEECFFFERLESNNYNTYRSRKYSDWYVALKRTGQYKLGSKTGPQKAILFLPMSAKS 288
.....160.....170.....180.....190.....200.....210.....220.....230.....240.....250.....260.....270.....280.....

```

