

TfoetusA1	<p>Sequence</p> <p>MSSSADRFRITIFSQEGRWQVEYAFKAVKQAEVTAIGAKGANSLLVVAVQKKVQDKLIDPSTVTHMFRIT EHVGACLIIGLLPDVLYICRRLRYDAAQFQFNGFEI PVNILLSRLSEIHQIESQYSGARPTGVSAIIFG FEPSKNDFFALYKVEPSGFTSGYRAVSCGVKEVEAMSALEKKYESFPTEKETAEFVISTLQTVVGLDFEA REVEVAVVTRENTAYRVMKADEVEKVLTAVAEKD</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_6 cd03754 264.10 8.83E-89 20S proteasome subunit alpha 1 (K02730)</p>
TfoetusA2	<p>Sequence</p> <p>MGDSDFSLTTTFSSGGKLGQIEHALKAVSLSGQCQVGVKAKDGAVIACESKPPSPLAERDTNLKVQAINPN VGCVYSGIITDYRVLLKLRKEAMKYKLRGVEVMPREVVKSAAAINQSYTQSGGVRPFGVSLLIIGWE DIPTLWQVDPSTFWAWKATALGKRSDGSRFTFLERRYNEVDDAIHTAISTLKEGFDGQLTKDLIEI GVVDKTHKFRTLSTEEIEDFLTEV</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_2 cd03750 324.66 1.85E-112 20S proteasome subunit alpha 2 (K02726)</p>
TfoetusA3	<p>Sequence</p> <p>MSYRYDAGTTTFSSDGRILQVEYAIQSINQAGTAIGVQFSNGVVLAAEKNTGRLVDYLFPEKMAKLDE HVITALAGMTADANNLVDVFMRSQAQSYLKYTGEPMPVEQLVRRVCDKHSYTQYGLRPGVGSFLIAGY DRHKGCLYLTDPSTGNGGKATAIGENNTAQSILKSSYKDDMNATEAMDLTVKVLCCTLSTSLSD KLEFVSVLQYSEKTPKVRILTAEVEHLMKRFEDTIKATTEEKE</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_4 cd03752 328.92 5.75E-114 20S proteasome subunit alpha 3 (K02728)</p>
TfoetusA4	<p>Sequence</p> <p>MSHYARSI TRFSPDGRLEFQIDHAQAQVQRGTTVVAVQSKDLIVIAVEKLTIAKLQDPCTFRKVVVA LDDHVMCAFAGLHADARVLVQKAQVEQCQSHRLTYEDPITVERIARHIATLQLKYTQSGGVRPFGV ATLVCGFDSTTKQPHIYETLPSGAYAEWKARTIGRHDQTVMEYLEKHYKDDMTDDDAIKMAVGS LEVVENGAKNLEVALMKCGEPMVLMKEQLAPLIEAINKK</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_7 cd03755 322.39 8.68E-112 20S proteasome subunit alpha 4 (K02731)</p>
TfoetusA5	<p>Sequence</p> <p>MFQSSSEYDRNVNTFSPDGRLLQVEYAIQAVKLGSSAIAILCPEGVVFVEKRLSSPLLISSVERVYA IDEHIGVVLAGYPADGRTMVDHMRVAAQDHRFSFGEPIGIRAVTESVCDLALSFGEGRRKREGQMSRPF GAALLVSGIDNGKPFLEHTDPSGTYTQCRARAIGGSEGAETILRDSYHDGMTLEEAENLALSTLRQVI QEKLSENNIEVACANVQTGRFKVYSAEQRAIVERLPPPLIQE</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_5 cd03753 331.22 5.56E-115 20S proteasome subunit alpha 5 (K02729)</p>
TfoetusA6	<p>Sequence</p> <p>MFRSRYDGDTTTFSPGRLLQVENAMKAVQQGMSTVGI RSTHAVIACIMHSPSEFSSFPKIFKIDEH IGVTISGLTADGRGLCKMLRSECLRHKFVYGTESKVVSTLADFIADRSQNKTKQVKGKRPYGVLLMIGAN PADGPRLFETCPGQNYEYDAQSIGRRSQAAKTYLEQNLPEFHNSTRDELIKHALKALFDCRAKEENGL ECFAVGVVGVDEPFMIIEGDALRPYFSE</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_1 cd03749 297.28 6.74E-102 20S proteasome subunit alpha 6 (K02725)</p>
TfoetusA7	<p>Sequence</p> <p>MSGAGSGYDFNAFTFSPDGRLEFQVEYATKAVDKAALGVRCSGVLFAVEKNLSSKLLTPGGNPRTFW IDTHIACATVGYRSDSYAAVLQARKEAANYFETFGCKITVPELVSRVAHTFHSSSHAFSSIRPYGCALLI GSLEGPSLYALEPNGQYFGYYACCFGKGSLLARAELQRTDWGEKTVEEAVPLVANI IKELHEAQNKTWE IEMLVWCQASEGKPKVPESLFVAAQ</p> <p>Motif Identifier Score E-value KEGG orthology Proteasome_alpha_type_3 cd03751 247.19 3.13E-82 20S proteasome subunit alpha 7 (K02727)</p>