

TfoetusB1	<p>Sequence</p> <p>MTNPNALVEMTGTTLFAIKTADGVVFAADSRSTAGSTVACRTSNKITQISSKVFVARTGSTADTQALAR YGHYILNVLTMNSESEVKRPVLVASQYLRKVLQSNKEYLSASLICGGIEDDGTPSLYEINTSGMAIPRD FCLNGSGSSYILAFCDKEYRDNMTMEEATKFAIEAVTNAIIRDGASGGFVNIVQVTKDQTKRFVVMPKD QPYNHSVVKT</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_6 cd03762 197.06 2.65E-63 20S proteasome subunit beta 1 (K02738)</p>
TfoetusB2	<p>Sequence</p> <p>MQNKGFDFDLIARNESLKVQPKKLKTGTTIAAAVFDGGVVLGADTRATAGPVVAVKDEFKIHYSNDNIW CCGAGTAADNDQITTLISSKLRLYTMNSGMQPRVYQAATLLTSRLFKYQGYIEAYLIVAGVDFQGPSVY TMYANGSCSKSPFSTMGSGSMAAVSVLEAGWKPKMSEEACKELVADAI EAGITNDLGSGSNVNL CVIKN TGHEYISHYRVTNERTFRMNAPIKRGTEIVISSSTTKPLHPPEVHLEILDGVAEEAAA</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_7 cd03763 292.18 8.63E-100 20S proteasome subunit beta 2 (K02739)</p>
TfoetusB3	<p>Sequence</p> <p>MSSGIDTQNGVSVVAMAGDHCVAIAADRRLGAQMOTVSTNFKKIFQINDYIHVGMGLATDIDTVYEKL RYDVNLLLELREERQLEPIRFMSLVRSLLEYHRFGPYFVSPVIAGLDPATNEPYLATSDSIGAFNQPKDF AVAGTSEESLYGICESMWRPGLNPDELFDVISQCLIAAVERDGLSGWGAVVHIITPESVVVKEIKTRMD</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_3 cd03759 317.65 1.23E-110 20S proteasome subunit beta 3 (K02735)</p>
TfoetusB4	<p>Sequence</p> <p>MLSIVGLCGPDWVLI AADSSVSSSIICMSEEDRIAEIGKHNALALAGETGDALQLSEYIIGNVALYKF INSVELTTDAISHYIRNEMAKAVRKNPYQVNMLLAGYDEKPSLYYLDYLGTRQKIPFGAQGYCAYFVLS VFDKFEENMSLEKGEVMKKALDQIKQRFTIAPHGFIVKLIDANGIQKINLD</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_2 cd03758 236.33 5.68E-79 20S proteasome subunit beta 4 (K02734)</p>
TfoetusB5	<p>Sequence</p> <p>MSSLQFPQHVDVIDGENQAPLQPANVEDPCGF IKNHLHFNYTNDAPDSNR IAAHYGTTCLSFIYKGGIVV AVDSRATGGSFIFSGTVMKILDIAPNMI GTMAGGAADCQYWLRLSRLSRLHNFYRQQPLTVAAASKIL VNELYSYKGYNLSIGTMICGYDSTGPHIFYIDNDGSRLEGKRFSVSGSTYAYGVLDTCYKYEMTKEEA CDLGRKAIYHATYRDSGSGGRVTVVHIDENGAQRISQTDVFDIHDFDKKVE</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_5 cd03761 313.80 2.04E-108 20S proteasome subunit beta 5 (K02737)</p>
TfoetusB6	<p>Sequence</p> <p>MHSEVAGKIEDKKGWSPYDNHGGTSVGI V GKDFVAIGTDTRLSSNYSISCRHKSRVFQMTSKAMIVAT GFDGDI DAFVTRIRQILVRYQOEHFKE MSTESLALCVSNILYSKRFFPYINILVGGIGLKDEGLLYGY DPVGTLECLNYDAHGTGSPMAMPILDNHFGSMHNTTPFPHPVEVDDAVNLIRDIMASVSERDIYTGDCCL QVAVMKSDGNLTIITEYELPAH</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Proteasome_beta_type_1 cd03757 214.04 2.28E-69 20S proteasome subunit beta 6 (K02732)</p>
TfoetusB7	<p>Sequence</p> <p>MSGYKKSQNPITTTNSIIAAKYKDGILLASDRAVSYGSCFKFANVSHFAKLT PNI IIGGTGELADFQE LVDVLRSIIVDEECKNNGESLTPSEVSNYIKRLMYERRSKMNPV MRCILAGIDKDGSKLLTATDLYGT QWEDEYVASGYGAHMQGVQIPKALKSADVTRDVMDA I KEVFI GLTARHSTMSGPIEFVDVTANGIEFL DPIEIVPNWDVLDDEDWAQ</p> <p>Motif Identifier Score E-value KEGG orthology</p> <p>Ntn_hydrolase super family cl00467 173.53 7.65E-54 20S proteasome subunit beta 7 (K02736)</p>