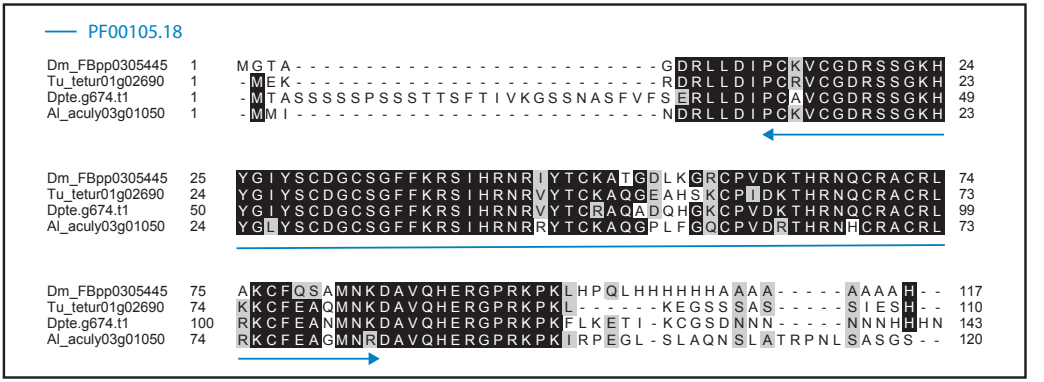
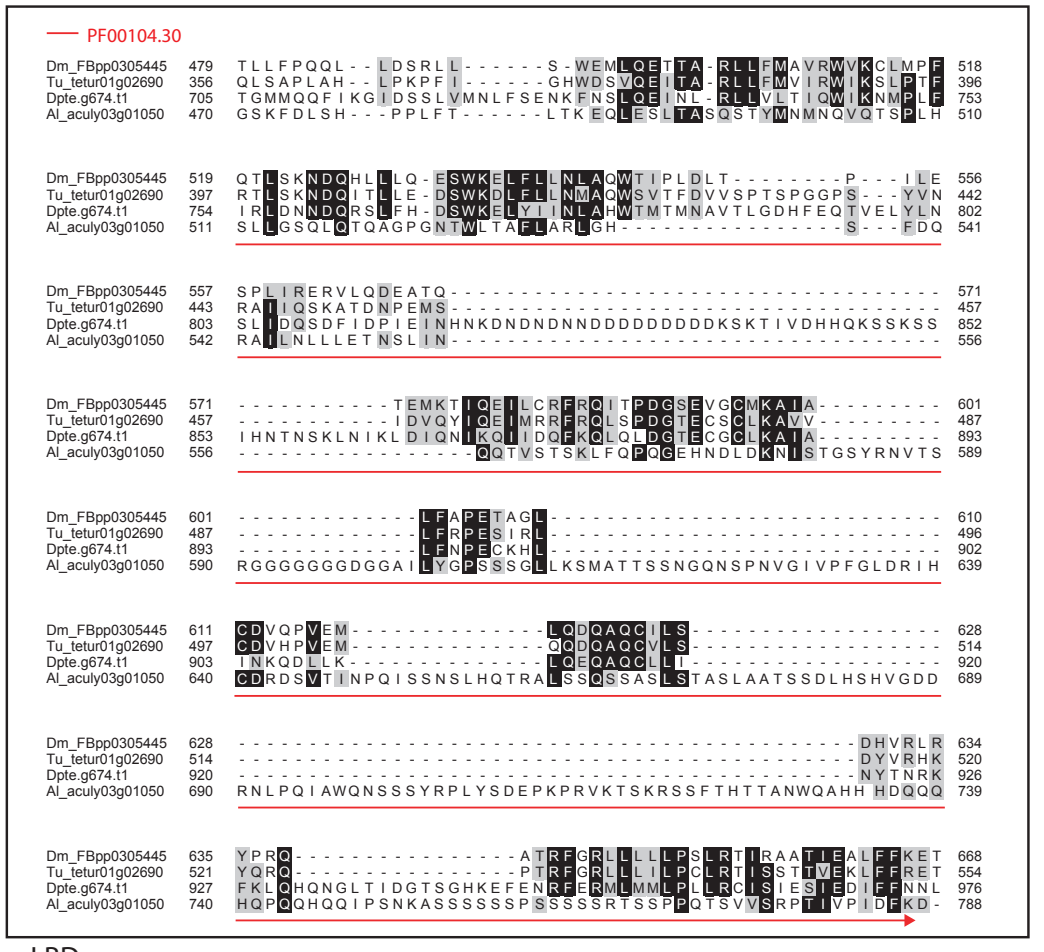

PF00105.18
Dm_FBpp0305445 1 MGTA-------------------------GDRLLDIPCKVCGDRSSGKH 24
Tu_tetur01g02690 1 -MEK-------------------------RDRLLDIPCRVCGDRSSGKH 23
Dpte.g674.t1 1 -MTASSSSSPSSSTTSFTIVKGSSNASFVFSERLLDIPCAVCGDRSSGKH 49
Al_aculy03g01050 1 -MMI-------------------------NDRLLDIPCKVCGDRSSGKH 23
Dm_FBpp0305445 25 YGIYSCDGCSGFFKRSIHRNRIYTCKATGDLKGRCPVDKTHRNQCRACRL 74
Tu_tetur01g02690 24 YGIYSCDGCSGFFKRSIHRNRVYTCKAQGEAHSKCPIDKTHRNQCRACRL 73
Dpte.g674.t1 50 YGIYSCDGCSGFFKRSIHRNRVYTCRAQADQHGKCPVDKTHRNQCRACRL 99
Al_aculy03g01050 24 YGLYSCDGCSGFFKRSIHRNRRYTCKAQGPLFGQCPVDRTHRNHCRACRL 73
Dm_FBpp0305445 75 AKCFQSAMNKDAVQHERGPRKPKLHPQLHHHHHHAAAA-----AAAAH-- 117
Tu_tetur01g02690 74 KKCFEAQMNKDAVQHERGPRKPKL------KEGSSSAS-----SIESH-- 110
Dpte.g674.t1 100 RKCFEANMNKDAVQHERGPRKPKFLKETI-KCGSDNNN-----NNNHHHN 143
Al_aculy03g01050 74 RKCFEAGMNRDAVQHERGPRKPKIRPEGL-SLAQNSLATRPNLSASGS-- 120

DBD


PF00104.30
Dm_FBpp0305445 479 TLLFPQQL--LDSRLL------S-WEMLQETTA-RLLFMAVRWVKCLMPF 518
Tu_tetur01g02690 356 QLSAPLAH--LPKPFI------GHWDSVQEITA-RLLFMVIRWIKSLPTF 396
Dpte.g674.t1 705 TGMMQQFIKGIDSSLVMNLFSENKFNSLQEINL-RLLVLTIQWIKNMPLF 753
Al_aculy03g01050 470 GSKFDLSH---PPLFT------LTKEQLESLTASQSTYMNMNQVQTSPLH 510
Dm_FBpp0305445 519 QTLSKNDQHLLLQ-ESWKELFLLNLAQWTIPLDLT--------P---ILE 556
Tu_tetur01g02690 397 RTLSKNDQITLLE-DSWKDLFLLNMAQWSVTFDVVSPTSPGGPS---YVN 442
Dpte.g674.t1 754 IRLDNNDQRSLFH-DSWKELYIINLAHWTMTMNAVTLGDHFEQTVELYLN 802
Al_aculy03g01050 511 SLLGSQLQTQAGPGNTWLTAFLARLGH----------------S---FDQ 541
Dm_FBpp0305445 557 SPLIRERVLQDEATQ---------------------------------- 571
Tu_tetur01g02690 443 RAIIQSKATDNPEMS---------------------------------- 457
Dpte.g674.t1 803 SLIDQSDFIDPIEINHNKDNDNDNNDDDDDDDDDKSKTIVDHHQKSSKSS 852
Al_aculy03g01050 542 RAILNLLLETNSLIN---------------------------------- 556
Dm_FBpp0305445 571 ----------TEMKTIQEILCRFRQITPDGSEVGCMKAIA--------- 601
Tu_tetur01g02690 457 ----------IDVQYIQEIMRRFRQLSPDGTECSCLKAVV--------- 487
Dpte.g674.t1 853 IHNTNSKLNIKLDIQNIKQIIDQFKQLQLDGTECGCLKAIA--------- 893
Al_aculy03g01050 556 ----------------QQTVSTSKLFQPQGEHNDLDKNISTGSYRNVTS 589
Dm_FBpp0305445 601 ----------LFAPETAGL---------------------------- 610
Tu_tetur01g02690 487 ----------LFRPESIRL---------------------------- 496
Dpte.g674.t1 893 ----------LFNPECKHL---------------------------- 902
Al_aculy03g01050 590 RGGGGGGGDGGAILYGPSSSGLLKSMATTSSNGQNSPNVGIVPFGLDRIH 639
Dm_FBpp0305445 611 CDVQPVEM----------LQDQAQCILS---------- 628
Tu_tetur01g02690 497 CDVHPVEM----------QQDQAQCVLS---------- 514
Dpte.g674.t1 903 INKQDLLK----------LQEQAQCLLI---------- 920
Al_aculy03g01050 640 CDRDSVTINPQISSNSLHQTRALSSQSSASLSTASLAATSSDLHSHVGDD 689
Dm_FBpp0305445 628 ----------DHVRLR 634
Tu_tetur01g02690 514 ----------DYVRHK 520
Dpte.g674.t1 920 ----------NYTNRK 926
Al_aculy03g01050 690 RNLPQIAWQNSSSYRPLYSDEPKPRVKTSKRSSFTHTTANWQAHH HDQQQ 739
Dm_FBpp0305445 635 YPRQ----------ATRFGRLLLLLPSLRTIRAATIEALFFKET 668
Tu_tetur01g02690 521 YQRQ----------PTRFGRLLLILPCLRTISSTTVEKLFFRET 554
Dpte.g674.t1 927 FKLQHQNGLTIDGTSGHKEFENRFERMLMMLPLLRCISIESIEDIFFNNL 976
Al_aculy03g01050 740 HQPQQHQQIPSNKASSSSSSPSSSSSRTSSPPQTSVVSRPTIVPIDFKD- 788

LBD