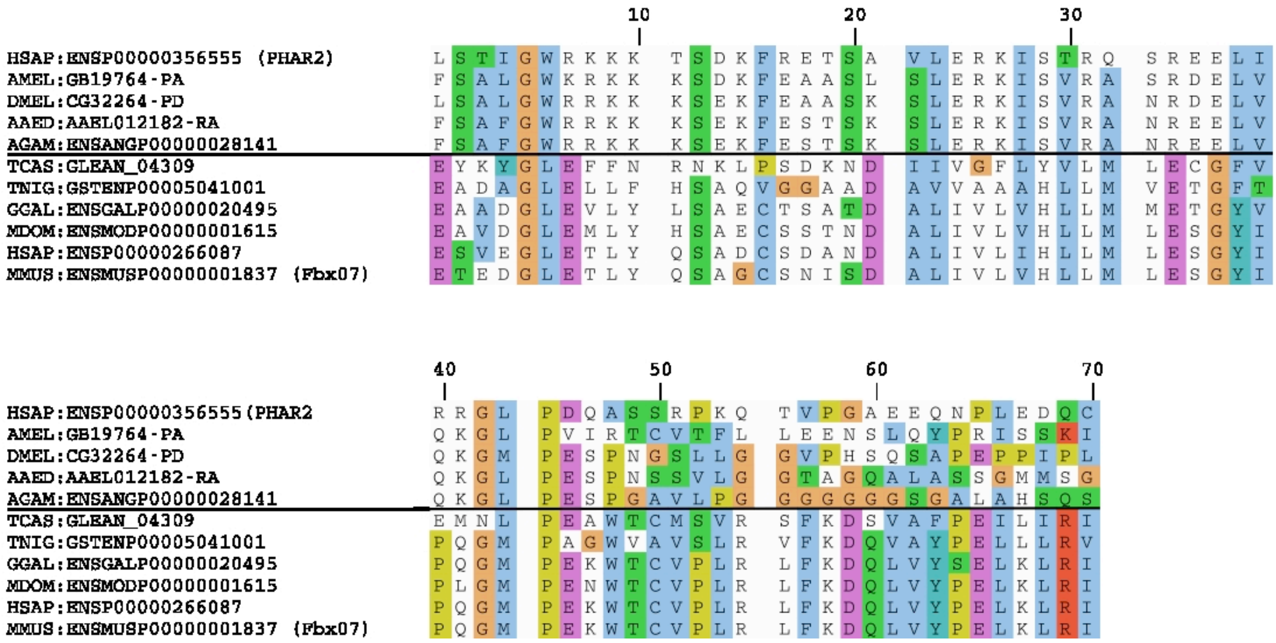

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                                          10                   20                   30
HSAP:ENSP00000356555 (PHAR2)      L S T I G W R K K K  T S D K F R E T S A  V L E R K I S T R Q  S R E E L I
AMEL:GB19764-PA                   F S A L G W K R K K  K S D K F E A A S L  S L E R K I S V R A  S R D E L V
DMEL:CG32264-PD                   L S A L G W R R K K  K S E K F E A A S K  S L E R K I S V R A  N R D E L V
AAED:AAEL012182-RA                F S A F G W R R K K  K S E K F E S T S K  S L E R K I S V R A  N R E E L V
AGAM:ENSANGP00000028141           F S A F G W R R K K  K S E K F E S T S K  S L E R K I S V R A  N R E E L V
TCAS:GLEAN_04309                  E Y K Y G L E F F N  R N K L P S D K N D  I I V G F L Y V L M  L E C G F V
TNIG:GSTENP00005041001            E A D A G L E L L F  H S A Q V G G A A D  A V V A A A H L L M  V E T G F T
GGAL:ENSGALP00000020495           E A A D G L E V L Y  L S A E C T S A T D  A L I V L V H L L M  M E T G Y V
MDOM:ENSMODP00000001615           E A V D G L E M L Y  H S A E C S S T N D  A L I V L V H L L M  L E S G Y I
HSAP:ENSP00000266087              E S V E G L E T L Y  Q S A D C S D A N D  A L I V L I H L L M  L E S G Y I
MMUS:ENSMUSP00000001837 (Fbx07)   E T E D G L E T L Y  Q S A G C S N I S D  A L I V L V H L L M  L E S G Y I

                                  40                   50                   60                   70
HSAP:ENSP00000356555(PHAR2        R R G L  P D Q A S S R P K Q  T V P G A E E Q N P L E D Q C
AMEL:GB19764-PA                   Q K G L  P V I R T C V T F L  L E E N S L Q Y P R I S S K I
DMEL:CG32264-PD                   Q K G M  P E S P N G S L L G  G V P H S Q S A P E P P I P L
AAED:AAEL012182-RA                Q K G L  P E S P N S S V L G  G T A G Q A L A S S G M M S G
AGAM:ENSANGP00000028141           Q K G L  P E S P G A V L P G  G G G G G G S G A L A H S Q S
TCAS:GLEAN_04309                  E M N L  P E A W T C M S V R  S F K D S V A F P E I L I R I
TNIG:GSTENP00005041001            P Q G M  P A G W V A V S L R  V F K D Q V A Y P E L L L R V
GGAL:ENSGALP00000020495           P Q G M  P E K W T C V P L R  L F K D Q L V Y S E L K L R I
MDOM:ENSMODP00000001615           P L G M  P E N W T C V P L R  L F K D Q L V Y P E L K L R I
HSAP:ENSP00000266087              P Q G M  P E K W T C V P L R  L F K D Q L V Y P E L K L R I
MMUS:ENSMUSP00000001837 (Fbx07)   P Q G M  P E K W T C V P L R  L F K D Q L V Y P E L K L R I
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Additional data file 7. Alignment of Fbox7 proteins (below the black line) and Phosphatase and actin regulators PHAR2 (above the black line). Tribolium Glean_04309 is more similar to vertebrate Fbox7 proteins than to insect proteins.