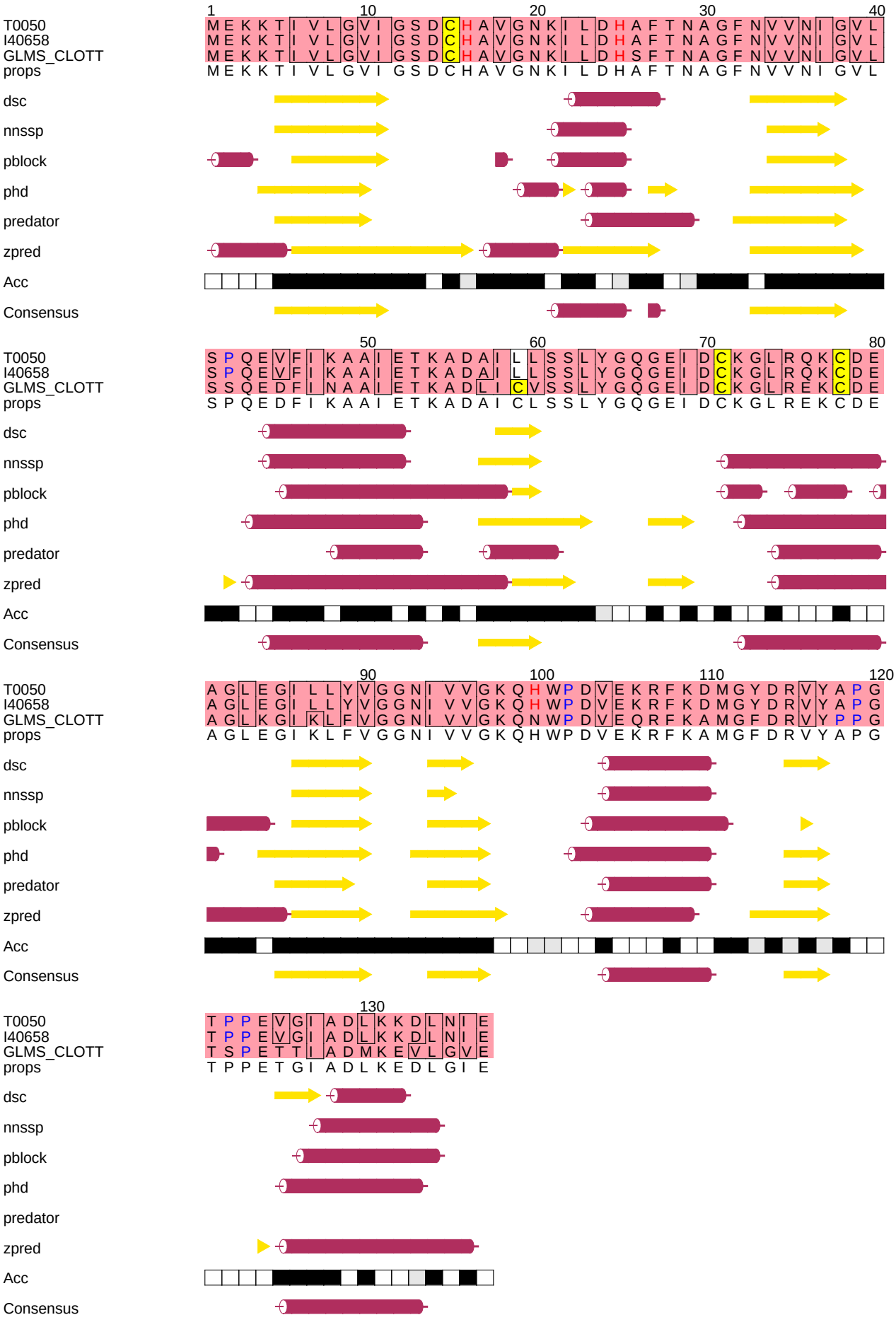
1 10 20 30 40
T0050 M E K K T I V L G V I G S D C H A V G N K I L D H A F T N A G F N V V N I G V L
I40658 M E K K T I V L G V I G S D C H A V G N K I L D H A F T N A G F N V V N I G V L
GLMS_CLOTT M E K K T I V L G V I G S D C H A V G N K I L D H S F T N A G F N V V N I G V L
props M E K K T I V L G V I G S D C H A V G N K I L D H A F T N A G F N V V N I G V L
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
50 60 70 80
T0050 S P Q E V F I K A A I E T K A D A I L L S S L Y G Q G E I D C K G L R Q K C D E
I40658 S P Q E V F I K A A I E T K A D A I L L S S L Y G Q G E I D C K G L R Q K C D E
GLMS_CLOTT S S Q E D F I N A A I E T K A D L I C V S S L Y G Q G E I D C K G L R E K C D E
props S P Q E D F I K A A I E T K A D A I C L S S L Y G Q G E I D C K G L R E K C D E
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
90 100 110 120
T0050 A G L E G I L L Y V G G N I V V G K Q H W P D V E K R F K D M G Y D R V Y A P G
I40658 A G L E G I L L Y V G G N I V V G K Q H W P D V E K R F K D M G Y D R V Y A P G
GLMS_CLOTT A G L K G I K L F V G G N I V V G K Q N W P D V E Q R F K A M G F D R V Y P P G
props A G L E G I K L F V G G N I V V G K Q H W P D V E K R F K A M G F D R V Y A P G
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
130
T0050 T P P E V G I A D L K K D L N I E
I40658 T P P E V G I A D L K K D L N I E
GLMS_CLOTT T S P E T T I A D M K E V L G V E
props T P P E T G I A D L K E D L G I E
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus