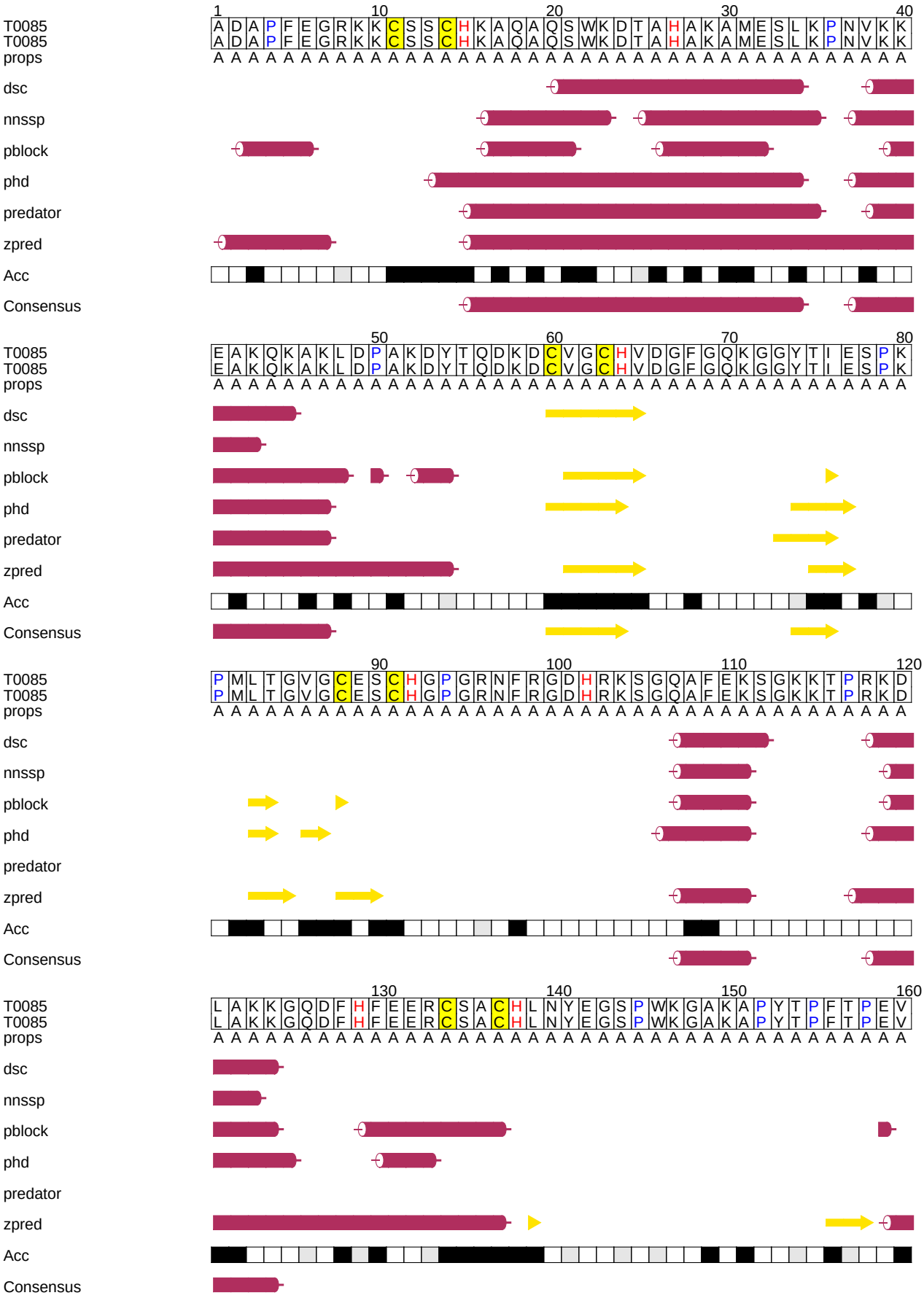
1 10 20 30 40
T0085 ADAPFEGRKKCSSSCHKAQAQSWKDTAHAKAMESLKPNVKK
T0085 ADAPFEGRKKCSSSCHKAQAQSWKDTAHAKAMESLKPNVKK
props AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
50 60 70 80
T0085 EAKQKAAKLLDDPAKDDYTQDDKDDCVGCHVDGFGQKGGGYTIESPK
T0085 EAKQKAAKLLDDPAKDDYTQDDKDDCVGCHVDGFGQKGGGYTIESPK
props AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
90 100 110 120
T0085 PMLTGVGCESSCHGPGRNFRGDDHRKSSGQAFEKSSGKKTTPRKD
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props AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
130 140 150 160
T0085 LAKKGQDDFHFEEERCSACCHLNYEGSPWKGAKAPYTPFTTPEEV
T0085 LAKKGQDDFHFEEERCSACCHLNYEGSPWKGAKAPYTPFTTPEEV
props AAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAAA
dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus

170
180
190
200
T0085
T0085
props
D A K Y T F K F D E M V K E V K A M H E H Y K L E G V F E G E P K F K F H D E F
D A K Y T F K F D E M V K E V K A M H E H Y K L E G V F E G E P K F K F H D E F
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dsc
nnssp
pblock
phd
predator
zpred
Acc
Consensus
210
T0085
T0085
props
Q A S A K P A K K G K
Q A S A K P A K K G K
A A A A A A A A A A A
dsc
nnssp
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phd
predator
zpred
Acc
Consensus