```
capb_chick  MSDQQLDCAL DLMRRLPPQQ IEKNLSDLID LVPSLCEDLL SSVDQPLKIA RDKVVGKDYL 60
capb_mouse  MSDQQLDCAL DLMRRLPPQQ IEKNLSDLID LVPSLCEDLL SSVDQPLKIA RDKVVGKDYL 60
capb_human  MSDQQLDCAL DLMRRLPPQQ IEKNLSDLID LVPSLCEDLL SSVDQPLKIA RDKVVGKDYL 60
capb_caeel  MGEQQLDCAL DLMRRLPPQH CDKNLTDLID LCPHLVDDLL STIDQPLKIA ADRETGKQYL 60
 Consensus  MSDQQLDCAL DLMRRLPPQQ IEKNLSDLID LVPSLCEDLL SSVDQPLKIA RDKVVGKDYL 60

capb_chick  LCDYNRDGDS YRSPWSNKYD PPLEDGAMPS ARLRKLEVEA NNAFDQYRDL YFEGGVSSVY 120
capb_mouse  LCDYNRDGDS YRSPWSNKYD PPLEDGAMPS ARLRKLEVEA NNAFDQYRDL YFEGGVSSVY 120
capb_human  LCDYNRDGDS YRSPWSNKYD PPLEDGAMPS ARLRKLEVEA NNAFDQYRDL YFEGGVSSVY 120
capb_caeel  LCDYNRDGDS YRSPWSNTYD PPLEDGQLPS EKRRKMEIEA NAAFESYRDL YFEGGVSSVY 120
 Consensus  LCDYNRDGDS YRSPWSNKYD PPLEDGAMPS ARLRKLEVEA NNAFDQYRDL YFEGGVSSVY 120

capb_chick  LWDLDH.GFA GVILIKKAGD GSKKIKGCWD SIHVVEVQEK SSGRTAHYKL TSTVMLWLQT 179
capb_mouse  LWDLDH.GFA GVILIKKAGD GSKKIKGCWD SIHVVEVQEK SSGRTAHYKL TSTVMLWLQT 179
capb_human  LWDLDH.GFA GVILIKKAGD GSKKIKGCWD SIHVVEVQEK SSGRTAHYKL TSTVMLWLQT 179
capb_caeel  FWDLDNGGFA GIVLIKKEGD GAKNITGCWD SIHVIEITER A..RQAHYKL TSTIMLWLQT 178
 Consensus  LWDLDH-GFA GVILIKKAGD GSKKIKGCWD SIHVVEVQEK SSGRTAHYKL TSTVMLWLQT 180

capb_chick  NKTGSGTMNL GGSLTRQMEK DETVSDSSPH IANIGRLVED MENKIRSTLN EIYFGKTKDI 239
capb_mouse  NKSGSGTMNL GGSLTRQMEK DETVSDCSPH IANIGRLVED MENKIRSTLN EIYFGKTKDI 239
capb_human  NKSGSGTMNL GGSLTRQMEK DETVSDCSPH IANIGRLVED MENKIRSTLN EIYFGKTKDI 239
capb_caeel  NKSSSGVMNL GGSLTRQHEM DAPINDQNTH LANMGRMIED QESKMRLTIN EIYFGKTKKV 238
 Consensus  NKSGSGTMNL GGSLTRQMEK DETVSDCSPH IANIGRLVED MENKIRSTLN EIYFGKTKDI 240

capb_chick  VNGLRSIDAI PDNQKYKQLQ RELSQVLTQR QIYIQPDN 277
capb_mouse  VNGLRSLDAI PDNHKFKQLQ RELSQVLTQR QVYIQPDN 277
capb_human  VNGLRSVQTF ADKSKQEALK NDLVEALKRK QQC~~~~~~ 272
capb_caeel  MSDLRSTEKQ SELEKQDEIV RELNNAMANR GN~~~~~~~ 270
 Consensus  VNGLRS-DAI PDN-KQKQLQ RELSQ-LTQR Q-YIQPD  278
```