```
logo                                          MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGV

NTDB id 1075710 ACJEH1 RS06810 WP 010947130.1 MTEKPQIINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGV   80
NTDB id 1275 LPP RS06810 WP 025519545.1       MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGV   80
NTDB id 1261 GCO85 RS07110 WP 025519545.1     MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGV   80
consensus                                     !!!!!!*!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!

logo                                          GIQFTGENSRYLCNKIETYLAGMLKSCQVTDTI

NTDB id 1075710 ACJEH1 RS06810 WP 010947130.1 GIQFTGENSRYLCNKIETYLAGMLKSCQVTDTI   113
NTDB id 1275 LPP RS06810 WP 025519545.1       GIQFTGENSRYLCNKIETYLAGMLKSCQVTDTI   113
NTDB id 1261 GCO85 RS07110 WP 025519545.1     GIQFTGENSRYLCNKIETYLAGMLKSCQVTDTI   113
consensus                                     !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved
X similar
X ≥ 50% conserved