```
logo                                        MTEKPQLINCSFPNEASLYMAYMPFVKGGGLFIRTQHVYSLGQSLELSVKLLNENEPYLVTGKIVWITPKGAQGNKPPGV

NTDB id 229551 CA929 RS07720 WP 027226909.1  MTEKPQLINCSFPNEASLYIAYMPFVKGGGLFIRTQHVYPLGQSLELSVKLLNENEPYLVAGKVVWITPKGAQGNKPPGV   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 229551 CA929 RS07720 WP 027226909.1  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 $\geq$ 50% conserved