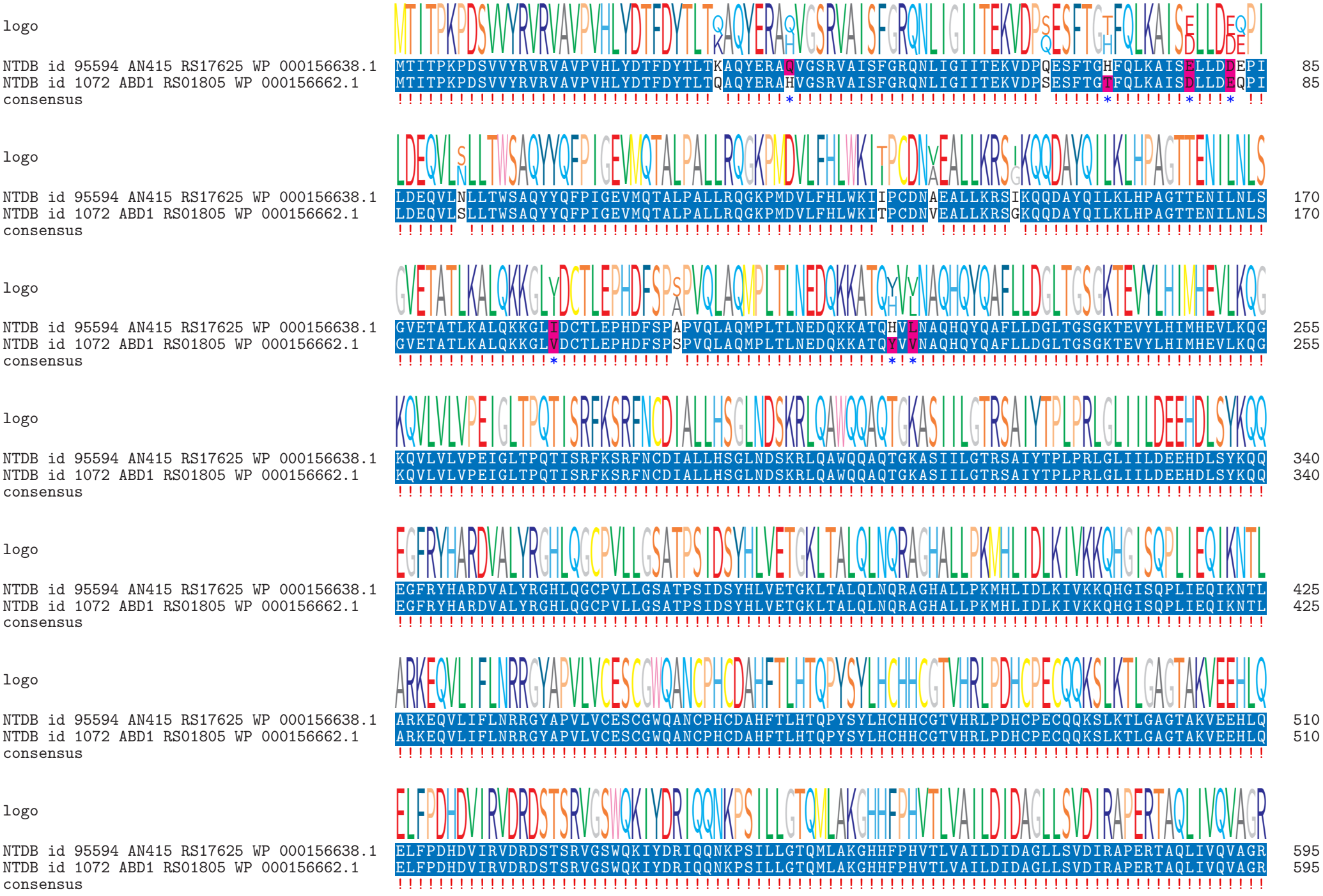

logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 MTITPKPDSVVYRVRVAVPVHLYDTFDYTLTKAQYERAQVGSRVAISFGRQNLIGIITEKVDPQESFTGHFQLKAISELLDDEPI 85
NTDB id 1072 ABD1 RS01805 WP 000156662.1 MTITPKPDSVVYRVRVAVPVHLYDTFDYTLTQAQYERAHVGSRVAISFGRQNLIGIITEKVDPSESFTGTFQLKAISDLLDFQPI 85
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 LDEQVLNLLTWSAQYYQFPIGEVMQTALPALLRQGKPMDVLFHLWKIIPCDNAEALLKRSIKQQDAYQILKLHPAGTTENILNLS 170
NTDB id 1072 ABD1 RS01805 WP 000156662.1 LDEQVLSLLTWSAQYYQFPIGEVMQTALPALLRQGKPMDVLFHLWKITPCDNVEALLKRSGKQQDAYQILKLHPAGTTENILNLS 170
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 GVETATLKALQKKGLIDCTLEPHDFSPAPVQLAQMPLTLNEDQKKATQHVLNAQHQYQAFLLDGLTGSGKTEVYLHIMHEVLKQG 255
NTDB id 1072 ABD1 RS01805 WP 000156662.1 GVETATLKALQKKGLVDCTLEPHDFSPSPVQLAQMPLTLNEDQKKATQYVVNAQHQYQAFLLDGLTGSGKTEVYLHIMHEVLKQG 255
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 KQVLVLVPEIGLTPQTISRFKSRFNCDIALLHSGLNDSKRLQAWQQAQTGKASIILGTRSAIYTPLPRLGLIILDEEHDLSYKQQ 340
NTDB id 1072 ABD1 RS01805 WP 000156662.1 KQVLVLVPEIGLTPQTISRFKSRFNCDIALLHSGLNDSKRLQAWQQAQTGKASIILGTRSAIYTPLPRLGLIILDEEHDLSYKQQ 340
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 EGFRYHARDVALYRGHLQGCPVLLGSATPSIDSYHLVETGKLTALQLNQRAGHALLPKMHLIDLKIVKKQHGISQPLIEQIKNTL 425
NTDB id 1072 ABD1 RS01805 WP 000156662.1 EGFRYHARDVALYRGHLQGCPVLLGSATPSIDSYHLVETGKLTALQLNQRAGHALLPKMHLIDLKIVKKQHGISQPLIEQIKNTL 425
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 ARKEQVLIFLNRRGYAPVLVCESCGWQANCPHCDAHFTLHTQPYSYLHCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ 510
NTDB id 1072 ABD1 RS01805 WP 000156662.1 ARKEQVLIFLNRRGYAPVLVCESCGWQANCPHCDAHFTLHTQPYSYLHCHHCGTVHRLPDHCPECQQKSLKTLGAGTAKVEEHLQ 510
consensus
logo
NTDB id 95594 AN415 RS17625 WP 000156638.1 ELFPDHDVIRVDRDSTSRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR 595
NTDB id 1072 ABD1 RS01805 WP 000156662.1 ELFPDHDVIRVDRDSTSRVGSWQKIYDRIQQNKPSILLGTQMLAKGHHFPHVTLVAILDIDAGLLSVDIRAPERTAQLIVQVAGR 595
consensus

logo 

```
NTDB id 95594 AN415 RS17625 WP 000156638.1  AGRGEHKGHVYLQTLRPDHPLLTTLIEKDYRAVAKQTLAERKVALLPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVD  680
NTDB id 1072 ABD1 RS01805 WP 000156662.1    AGRGEHKGHVYLQTLRPDHPLLTTLIEKDYRAVAKQTLAERKVALLPPYRYAVLIRAESKDRDYTLHFLNEAAEQLRQIAGDIVD  680
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

logo 

```
NTDB id 95594 AN415 RS17625 WP 000156638.1  IWGPIPAPMERKAGRYRAHMVILSADRARLHFYLRQWWAQLVHAPRQHQLRLSIDVDPQEFS  742
NTDB id 1072 ABD1 RS01805 WP 000156662.1    IWGPIPAPMERKAGRYRAHMVILSADRARLHFYLRQWWAQLVHAPRQHQLRLSIDVDPQEFS  742
consensus                                   !!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!!
```

X non conserved

X similar

X ≥ 50% conserved