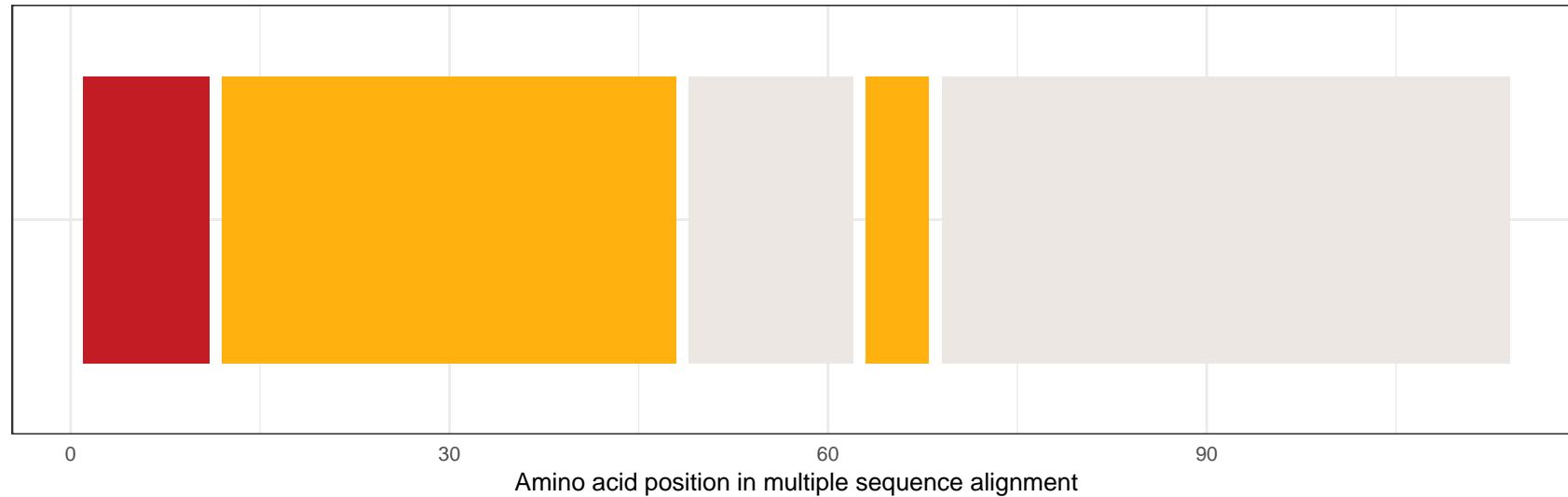


Protein topologies

Isoform ID (ENST)

ENST00000513264



Amino acid position in multiple sequence alignment

Protein topology

- Signal peptide
- Outside
- Trans membrane
- Inside
- Alignment gap