

# Protein topologies

Isoform ID (ENST)

ENST00000582480

ENST00000578509

ENST00000584284

ENST00000583376

ENST00000312648

0

50

100

150

200

250

Amino acid position in multiple sequence alignment

## Protein topology

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

