

Protein topologies

Isoform ID (ENST)

ENST00000595113
ENST00000405816
ENST00000398599
ENST00000221992
ENST00000617332
ENST00000615021
ENST00000595403

0

200

400

600

Amino acid position in multiple sequence alignment

Protein topology

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

