

# Protein topologies

Isoform ID (ENST)

ENST00000561896  
ENST00000566269  
ENST00000563941  
ENST00000566549  
ENST00000545450  
ENST00000382862  
ENST00000569566  
ENST00000563651

0

200

400

600

Amino acid position in multiple sequence alignment

## Protein topology

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

