

Protein topologies

Isoform ID (ENST)

ENST00000269571
ENST00000578502
ENST00000541774
ENST00000406381
ENST00000584601
ENST00000582818
ENST00000580074
ENST00000445658
ENST00000584450
ENST00000578709
ENST00000578199

0

500

1000

1500

Amino acid position in multiple sequence alignment

Protein topology

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

