
offset
Homo sapiens
1
cleavage sites - trypsin
modified residue
phospho.elm
pfam
secondary structure
homology model
peptide
chain
region of interest
metal ion-binding site
site
cross-link
sequence variant
splice variant
mobiDB
elm regex
signal peptide
Fibrinopeptide A
Fibrinogen alpha chain
Fib_alpha (6.9e-40)
Fibrinogen_aC (6.1e-27)
Fibrinogen_C (2.4e-71)
Disordered
tryps
P
Clea
Alpha-