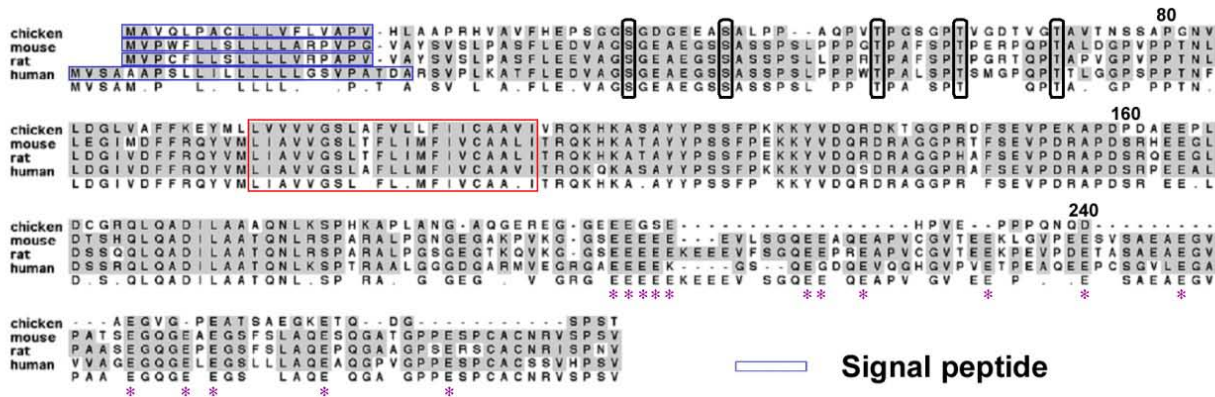




Identity (mouse / human) : 74.6%

-  Signal peptide
-  Transmembrane domain
-  Conserved glutamic acid residues
-  Potential O-glycosylation sites