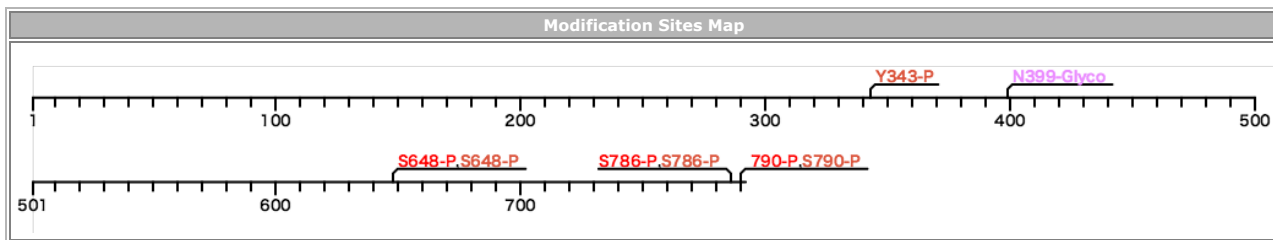


ID	Accession	GeneName	Chr.No.	Description
CADH6_HUMAN	P55285	CDH6	5p13.3 31193857..31329253	Cadherin-6



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
648	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
648	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
648	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
648	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
648	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
648	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
648	S	P	Paper	Mol Cell Proteomics 2009, 8(12), 2796-2808
648	S	P	Paper	Mol Cell Proteomics 2010, 9(6), 1047-1062

Protein Sequence	
MRTYRYFLLL FWVGGPYPTL STPLSKRTSG FPAKKRALEL SGNSKNELNR SKRSWMWNQF FLLEEYTGSD YQYVGKLHSD QDRGDGSLKY ILSGDGAGDL FIINENTGDI QA TKRLDREE KPVYILRAQA INRRTGRPVE PESEFIIKIH DINDNEPIFT KEVYTATVPE MSDVGTFFVQ VTATDADDPT YGNSAKVVYS ILQGQPYFSV ESETGIKTA LLNMDRE NRE QYQVVIQAKD MGGQMGGLSG TTTVNITLTD VNDNPPRFPQ STYQFKTPES SPPGTPIGRI KASDADVGEN AEIEYSITDG EGLDMFDVIT DQETQEGIIT VKKLLDFEKK KVVTLKVEAS NPYVEPRFLY LGPFKDSATV RIVVEDVDEP PVFSKLAYIL QIREDAQINT TIGSVTAQDP DAARNPVKYS VDRHTMDRI FNIDSGNGSI FTSKLLDRET LLWH NITVIA TEINNPQSS RVPLYIKVLD VNDNAPEFAE FYETFVCEKA KADQLIQLTL AVDKDDPYSG HQFSFSLAPE AASGSNFTIQ DNKDNTAGIL TRKNGYNRHE MSTYLLPV VI SDNDYPVQSS TGTVTVRVCA CDHHGNMQSC HAEALIHPTG LSTGALVAIL LCIVILLTV VLFAALRRQR KKEPLIISKE DIRDNIVSYN DEGGGEEDTQ AFDIGTLRNP EA IEDNKLRD DIVPEALFLP RRTPTARDNT DVRDFINQRL KENDTDPTAP PYDSLATYAY EGTGSVADSL SSLESVTTDA DQDYDYLSDW GPRFKKLADM YGGVDSDKD	

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing