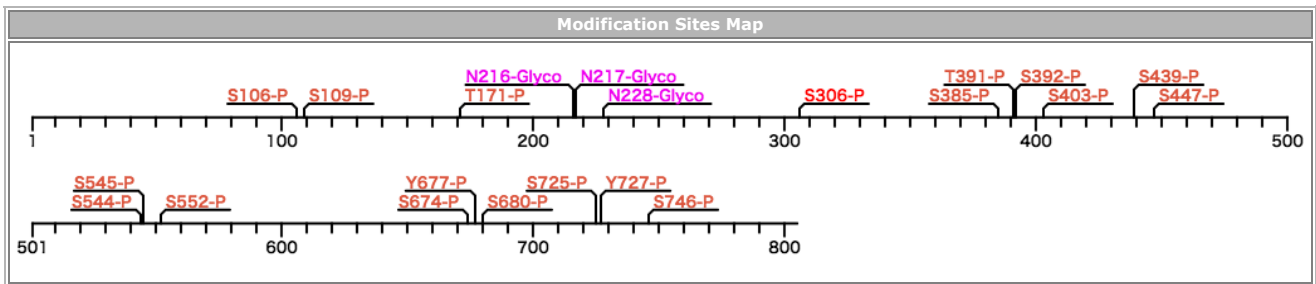


ID	Accession	GeneName	Chr.No.		Description
ENPL_HUMAN	P14625	HSP90B1	12q23.3	104323885..104347423	Endoplasmic



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
217	N	Glyco	Lab	130626_Gly_S_MGFPeaklist.mgf[F015371]
217	N	Glyco	Lab	130626_Gly_S_MGFPeaklist.mgf[F015371]
217	N	Glyco	Lab	130703_Gly_S_re_MGFPeaklist.mgf[F015387]
217	N	Glyco	Lab	130703_Gly_Y_re_MGFPeaklist.mgf[F015388]
217	N	Glyco	Lab	130703_Gly_Y_re_MGFPeaklist.mgf[F015388]

Protein Sequence	
MRALWVLGLC CVLLTFGSVR ADDEVDVDGT VEEDLGKSRE GSRTDDEVVQ REEEAIQLDG LNASQIRELR EKSEKFAFQA EVNRMMKLI NSLYKNKEIF LRELIS <sup>S</sup> NA <sup>S</sup> D ALD KIRLISL TDENALSGNE ELTVKIKCDK EKNLLHVTDT GVGMTREELV KNLGTIAKSG <sup>T</sup> SEFLNKMTE AQEDGQSTSE LIGQFGVGFY SAFLVADKVI VTSKH <sup>N</sup> NDTQ HIWESD S <sup>N</sup> EF SVIADPRGNT LGRGTTITLV LKEEASDYLE LDTIKNLVKK YSQFINFPIY VWSSKTETVE EPMEEEEAAK EEEKE <sup>S</sup> DDEA AVEEEEEEEK PKTKKVEKTV WDWELMNDIK PIWQRPSKEV EEDEYKAFYK SFSKESDDPM AYIHFTAEGE VTFK <sup>S</sup> ILFVP <sup>T</sup> SAPRGLFDE YG <sup>S</sup> KKSDYIK LYVRRVFITD DFHDMMPKYL NFVKGWV <sup>S</sup> D DLPLNV <sup>S</sup> RET LQQ HKLLKVI RKKLVRKTLD MIKKIADDKY NDTFWKEFGT NIKLGVIEDH SNRTRLAKLL RFQSSHHPTD ITSLDQYVER MKEKQDKIYF MAG <sup>S</sup> SRKEAE <sup>S</sup> SPFVERLLK KGYEVIY LTE PVDEYCIQAL PEFDGKRFQN VAKEGVKFDE SEKTKESREA VEKEFEPLLN WMKDKALKDK IEKAVVSQRL TESPALVAS QYGWSGNMR IMKAQAYQTG KDI <sup>S</sup> T <sup>N</sup> Y <sup>Y</sup> YA <sup>S</sup> QKKTFEINPR HPLIRDMLRR IKEDEDDKTV LDLAVLLET ATLR <sup>S</sup> GY <sup>L</sup> LP DTKAYGDRIE RMLRL <sup>S</sup> LNID PDAKVEEPEE EEPETAEDT TEDTEQDEDE EMDVGTDEEE ETA KESTAEK DEL	

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