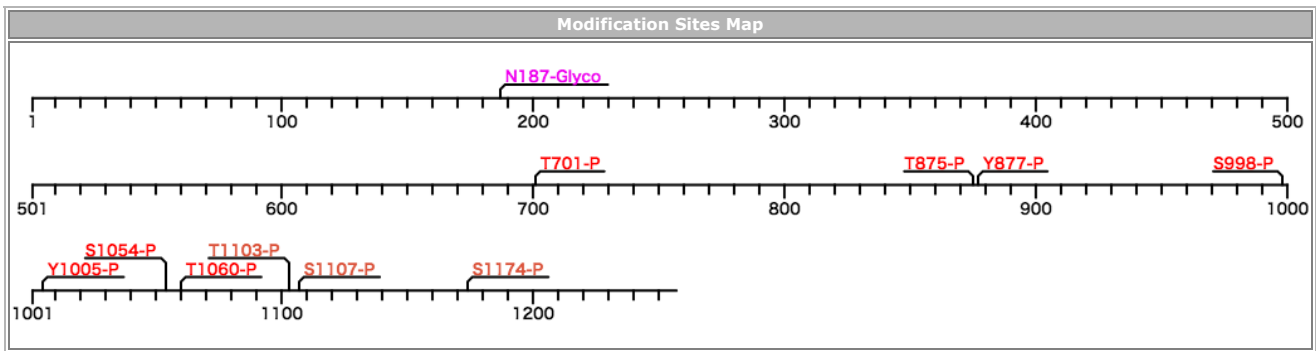


ID	Accession	GeneName	Chr.No.		Description
ERBB2_HUMAN	P04626	ERBB2	17q12	37844167..37886679	Receptor tyrosine-protein kinase erbB-2



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
187	N	Glyco	Lab	130626_Gly_S_MGFPeaklist.mgf[F015371]
187	N	Glyco	Lab	130626_Gly_Y_MGFPeaklist.mgf[F015373]
187	N	Glyco	Lab	130703_Gly_S_re_MGFPeaklist.mgf[F015387]

Protein Sequence	
MELAALCRWG LLLALLPPGA ASTQVCTGTD MKLRLPASPE THLDMLRHLY QGCQVVQGNL ELTYLPTNAS LSFQDIQEV QGYVLIHNPQ VRQVPLQRLR IVRGTQLFED NYA LAVLDNG DPLNNTTPTV GASPGGLREL QLRSLTEILK GGVLIQRNPQ LCYQDTILWK DIFHKNNQLA LTLIDTNRSR ACHPCSPMCK GSRCWGESSE DCQSLTRTVC AGGCA RCKGP LPTDCCHEQC AAGCTGPKHS DCLACLHFNH SGICELHCPA LVTYNTDTFE SMPNPEGRTY FGASCVTACP YNYLSTDVGS CTLVCPHNPQ EVTAEDGTQR CEKCSKP CAR VCYGLGMEHL REVRAVTSAN IQEFAGCKKI FGSLAFLPES FDGPASNTA PLQPEQLQVF ETLEEITGYL YISAWPDSLPL DLSVFNQLQV IRGRILHNGA YSLTLQGLGI S WLGLRSLRE LGSGLALIIH NTHLCFVHTV PWDQLFRNPH QALLHTANRP EDECVGEGLA CHQLCARGHC WPGPPTQCVN CSQFLRGQEC VEECRVLQGL PREYVNRHC L PCHPECQPQ NGSVTCFGPE ADQCACAHY KDPPFCVARC PSGVKPDL SY MPIWKFPDEE GACQPCPINC THSCVDLDDK GCPAEQRASP LTSIISAVVG ILLVVVLGVV FGIL IKRRQQ KIRKYTMRRLL QETELVEPL TPGSAMPNQA QMRILKETEL RKVKVLGSGA FGTVYKGIWI PDGENVKIPV AIKVLRENTS PKANKEILDE AYVMAGVGGSP YVSRLLGI CL TSTVQLVTQL MPYGCLLDHV RENRGRGSGQ DLLNWCMQIA KGMSYLEDVR LVHRDLAARN VLVKSPNHVK ITDFGLARLL DIDEYHAD GGKVPKWKMA LESILRRRFT HQSDVWSYGV TVWELMTFGA KPYDGIPARE IPDLEKGER LPQPICTID VYMIMVKCWM IDSECRPRFR ELVSEFSRMA RDPQRFVVIQ NEDLGPA S PL DSTFYRSLL DD DMGDLDVA EEYLVPPQGF FCPDPAPGAG GMVHHRHRSSTR SGGGDLT LGLEPSEEEA PRSPLAPSEG AGSDVFDGDL GMGAAGLQS LPTHDP S PLQ RYSEDPVPL PSE TDGYVAP LTCSPQPEYV NQPDVVRPQPP SPREGPLPAA RPAGATLERP KTLSPGKNGV VKDVFAGGA VENPEYLTPQ GGAAPQPHPP PAFSPAFDNL YYWDQDPPER GAPPS TFKGT PTAENPEYLG LDVPV	

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