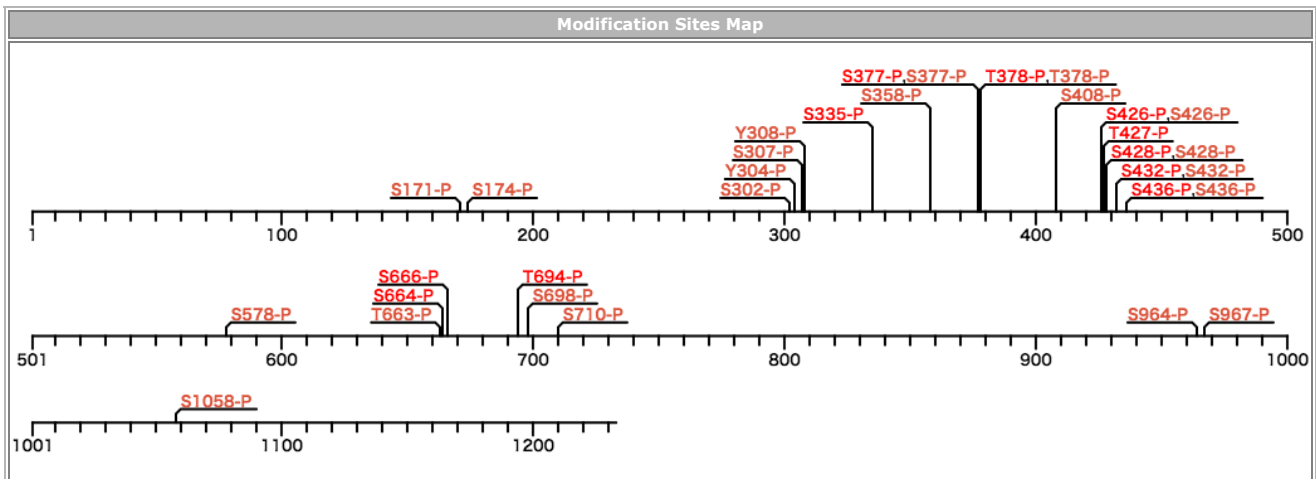


ID	Accession	GeneName	Chr.No.	Description	
EHBP1_HUMAN	Q8NDI1	EHBP1	2p15	62900986..63273622	EH domain-binding protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
377	S	P	Lab	100520-GIST-IM1.mgf[F017509]
377	S	P	Lab	100520-GIST-IM3.mgf[F017514]
377	S	P	Lab	100520-GIST-R3.mgf[F017519]
377	S	P	Paper	Sci Signal 2009, 2(84), ra46
377	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MASVWKRLQR VGKHASKQF VASYQELMVE CTKKWQPKL VVVWTRRSRR KSKAHSWQP GIKNPYRGVV VWPVPENIEI TVTLFKDPHA EEFEDKEWTF VIENESPSGR RKALATSSIN MKQYASPMPT QTDVCLKFKP LSKKVSAAL QFSLSCIFLR EGKATDEDMQ SLASLMSMKQ ADIGNLDDFE EDNEDDENR VNQEKAANKI TEIVNQLNAL SS LDEDQDDC IKQANMRSK SASSSEELIN KLNFLDEAEK DLATVNSNPF DDPDAAELNP FGDPSDEEPI TETASPRKTE DSFYNNSYNP FKEVQTPQYL NPFDEPEAFV TIKD SPQST KRKNIRPVD M SKLYADSSK TEEELDES N PFYEPKSTPP PNNLVNPVQE LETERRVKRK APAPPVLS PK TGVLNENTVS AGKDLSTSPK PPSIPSPVLG RKNPASQSLL VWCKEVTKNY RGVKITNFTT SWRNGLSFCA ILHHFRPDLI DYKSLNPQDI KENKKAYDG FASIGISRL EPDMLVLLAI PDKLTVMTYL YQIRAHFSGQ ELNVVQIEEN SSKS TYKVG N YETDTNSVD QEKFYAELSD LKREPELQQP ISGAVDFLSQ DDSVFVND SG VGESESEHQT PDDHLS PSTA SPYCRRTKSD TEPQKQSS GR TSGSDDPG ICSNT DSTQA QVLLGKRLLL KAEI LELSDL YVSDKKKDM S PPFICEETDE QKLQTLDIGS NLEKEKLENS RSLECRSDPE SPIKKTSLSP TSKLGYYSR DLDLAKKKHA SLRQTESDP D ADRTTLNHAD HSSKIVQHRL LSRQEELKER ARVLEQARR DAALKAGNKH NTNTATPFCN RQLSDQQDEE RRRQLRERAR QLIAEARSGV KMSELPSYGE MAAEKLKERS KASGDENDNI EIDTNEE IPE GFVVG GDEL TNLENDL DTP EQNSKLVLDL LKKLLEVQPQ VANSPSSAAQ KAVTESSEQD MKSGTEDLRT ERLQKTERF RNPVFSKDS TV RKTQLQSF SQYIENRPEM KRQRSIQEDT KKGNEEKA AI TETQRKPS ED EVLNKGFKDT SQYVVGELAA LENEQKQIDT RAALVEKRLR YLMDTGRNTE EEEAMMQEWF MLV NKKNALI RRMNQLSLL EKEHDLERRY E LLNREL RAML AIEDWQKTEA QKRREQLLLD ELVALVNKRD ALVRDLDAQE KQAEEDDEHL ERTLEQNKGK MAKKEEKCVL Q

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