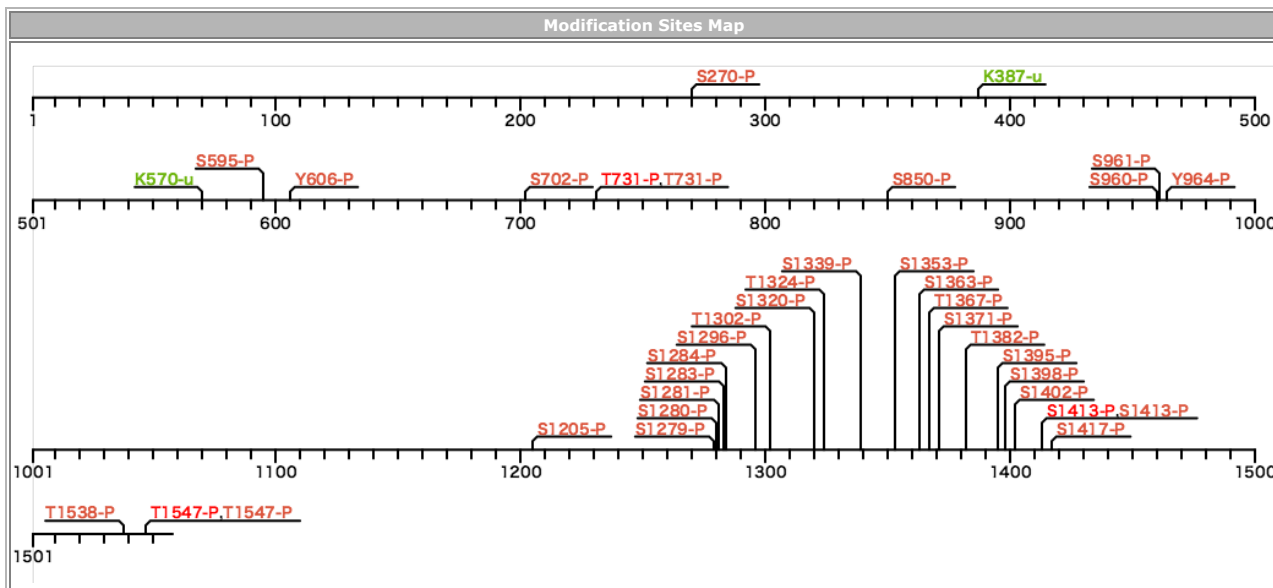


ID	Accession	GeneName	Chr.No.	Description
BAZ1A_HUMAN	Q9NRL2	BAZ1A	14q13.1 35221937..35344853	Bromodomain adjacent to zinc finger domain protein 1A



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
1547	T	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
1547	T	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
1547	T	P	Paper	Cell Rep 2014, 8(5), 1583-1594
1547	T	P	Paper	J Proteome Res 2013, 12(1), 260-271

Protein Sequence

MPLLHRKPFV RQKPPADLRP DEEVFYCKVT NEIFRHYDDF FERTILCNLS VWSCAVTGRP GLTYQEALQS EKARQNLQS FPELIIPVL YLTSLTHRSR LHEICDDIFA VVKDHY FVEE TVEVIRNNGA RLQCRILEVL PPSHQNGFAN GHVNSVDGET IIIISDSDSE TQSCSFQNGK KKDAIDPLLF KYKVQPTKKE LHESAIVKAT QISRRKHLFS RDKLKLFLKQ HCEPQDGVIK IKASSLSTYK IAEQDFSYFF PDDPPTFIF S PANRRRGRPP KRIHISQEDN VANKQTLASY RSKATKERDK LLKQEEMKSL AFEKAKLKRE KADALEAKK EKED KEKKRE ELKKIVEEER LKKKEEKERL KVEREKEREK LREEKR K YVE YLKQWSKPRE DMCCDDLKEL PEPTPVKTRL PPEIFGDALM VLEFLNAFGE LFDLQDEFPD GVTLEVLEE A LVGNDSGEP LCELLFFLTA IFQAIIEEEE EVAKEQLTDA DTKDLTEALD EDADPTKSAL SAVASLAAAW PQLHQGCSLK SLDLDSCTL S EILRLHILAS GADVTSANA K YRY QKRGGF D ATDDACMELR LSNP S LVKKL SSTS V YDLTP GEKMKILHAL CGKLLTLVST RDFIEDYVDI LRQAKQEFRE LKAEQHRKER EEAARIRKR KEEKLEKEQ KMKKEQ EKLK EDEQRNSTAD I S IGEERED FDTSES KDT EQKELDQDMV T EDEDDPGSH KRGRGRGRGQ NGFKEFTRQE QINCVTREP L TADEEEALKQ EHQRKEKELL EKIQSAIA CT NIFPLGRDRM YRRYWI FPSI PGLFIEEDYS GLTEDMLLPR PSSFQNNVQ S QDPQVSTKTG EPLMSESTN IDQGPRDHSV QLPKPVHKPN RWC FYSSCEQ LDQLIEALNS RGHRESALKE TLLQEKSRI C AQLARFSEEK FHFSDKPPQD SKPTYSRGR S S NAYDPSQMC AEKQLELRLR DFLLDIEDRI YQGTGAIKV TDRHIWRSAL ESGRYELLE ENKE NGI IKT VNEDVEEME I DEQTKVIVKD RLLGIKTETP STVSTNASTP QSVSSVVHYL AMALFQIEQG IERRFLKAPL DASDSGRSYK TVLDRWRESL LSSASLSQVF LHLSTLDR SV IWSKSILNAR CKICRKKGDA ENMVLCDGCD RGHHTYCVRP KLKTVPEDGW FCPECRPKQR SRRL S SRQP SLESDEDVED SMGGEDDEV D GDEEEGQSEE EEEYEVEQ DED DSQEEEEVSL PKRGRPVRL PVKTRGKLS S SFSSRGQQE PGRYP SRSQ S TPKTTVSSK TGRSLRKIN S APP TETKSLR IASRSTRHS H GPLQADVFE LL S PRRKRR G RKS ANNT PEN S PNFPNFRVI AT K SSEQSRS VNIA SKLS LQ ESEKRRCRK RQ S PEP S PVT LGRSSGRQG GVHELSAFEQ LVVELVRHDD SWPFLKLVSK IQVPDYDII K KPIALNIIR EKNVNC EYKL ASEFIDDIEL MFSNCFEYNP RNTSEAKAGT RLQAFFHIQA QKLG LHV T PS NVDQVST PPA AKKS R

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