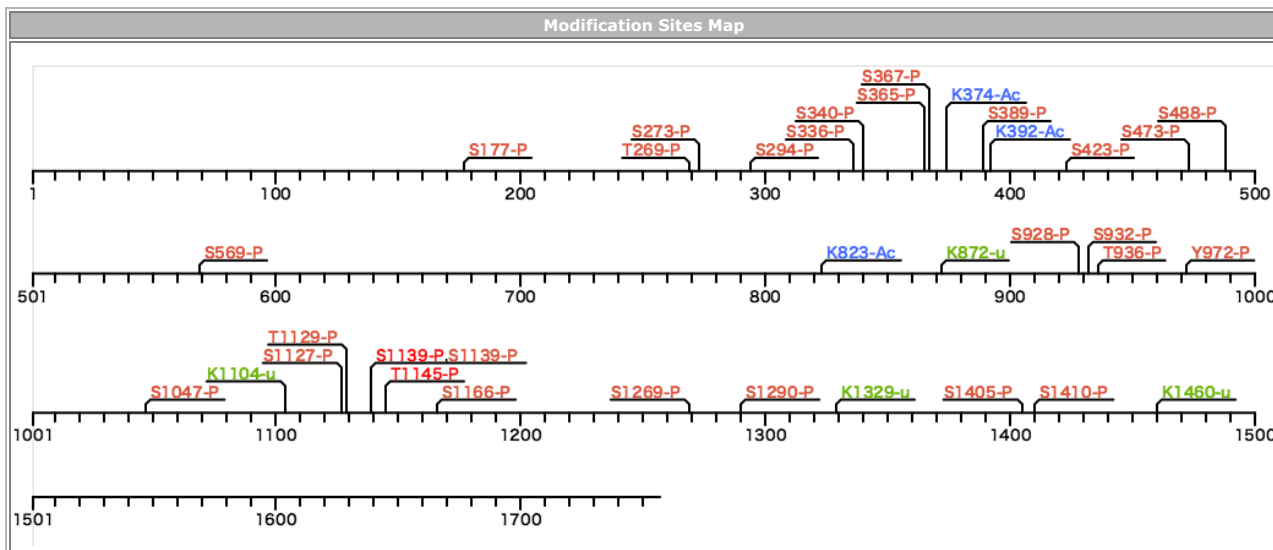


ID	Accession	GeneName	Chr.No.	Description
BCOR_HUMAN	Q6W2J9	BCOR	Xp11.4 39909068..40036582	BCL-6 corepressor



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
1139	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
1139	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
1139	S	P	Paper	J Proteomics 2014, 96, 253-262
1139	S	P	Paper	Sci Signal 2009, 2(84), ra46

**Protein Sequence**

MLSATPLYGN VHSWMNSERV RMCGASEDRK ILVNDGDASK ARLELREENP LNHNVVDAST AHRIDGLAAL SMDRTGLIRE GLRVPGNIVY SSLCGLGSEK GREAAATSLG G  
 LGFSSERNP EMQFKPNTPE TVEASAVSGK PPNGFSAIYK TPGGIQKSAV ATAEALGLDR PASDKQ<sup>S</sup>PLN INGASYLRLP WVNPYMEGAT PAIYPLDSP NKYSLNMYKA LLPQQ  
 SYSLA QPLYSVCTN GERFLYLPPP HYVGHIPSS LASPMLRS<sup>T</sup>P SA<sup>S</sup>PAIPPLV HCADKSLPWK MGV<sup>S</sup>SPGNPVD SHAYPHIQNS KQPRVPSAKA VTSGLPGDTA LLLPP<sup>S</sup>PRP  
<sup>S</sup>PRVHLPTQPA ADTYSEFHKK YARIS<sup>T</sup>SPSV ALS<sup>K</sup>PYMTVS SEFPAARLS<sup>N</sup>G<sup>K</sup>YPKAPEGG EGAQVPVGH<sup>A</sup>RKTAVQDRKD GS<sup>S</sup>PPLEKQ TVTKDVTDKP LDLSSKVVDV D  
 ASKADHMKK MAPTVLVHSR AG<sup>S</sup>GLVLVSGS EIPKETLS<sup>P</sup>PP GNGCAIYRSE IISTAPSSWV VPGPSPNEEN NGKSMSLKNK ALDWAIQQR SSSCPRMGGT DAVITNVSGS VS  
 SAGRPA<sup>S</sup>SA SPANANADG TKTSRSSVET TPSVIQHVQG PPATPAKHSS STSSKGAKAS NPEPSFKANE NGLPPSSIFL SPNEAFRSP<sup>P</sup>IPYPRSYL<sup>P</sup>Y PAPEGIAVSP LSLHGK  
 GPVY PHPVLLPNS LFPGLHAPK GLPYGLPTGR PEFVTYQDAL GLGMVHPMLI PHTPIETKE EKPERRSRSH ERARYEDPTL RNRFS<sup>E</sup>ILET SSTKLHPDVP TDKNLKPNPN  
 WNQGKTVVKS DKLVYVDLLR EEPDAKDTN VS<sup>K</sup>PSFAAES VGQSAEPPK<sup>P</sup>SVEPALQQHR DFIALREELG RISDFHETY<sup>T</sup>FK<sup>Q</sup>PVFTVSK DSVLAGTNKE NLGLPVSTPF LEP  
 PLGSDGP AVTFGKTQED PKPFCVGS<sup>A</sup>P PSVDV<sup>T</sup>PTYT KDGADAEASN DGKVLKPKPS KLAKRIANSA G<sup>V</sup>VGDRFKCV TTELYADSSQ LSREQRALQM EGLQEDSILC LPAAY  
 CERAM MRFSELEMKE REGGHPATKD SEMCKFS<sup>P</sup>AD WERLKG<sup>N</sup>QDK KPKSVTLEEA IAEQNESERC EYSVGNKHRD PFEAPEDKDL PVE<sup>K</sup>YFVERQ PVSEPPADQV ASDMP  
 HS<sup>P</sup>T<sup>L</sup>RVDRKRKVS<sup>G</sup>DSSHT<sup>T</sup>ETTA<sup>E</sup>EVPEPDL<sup>L</sup>KA KRRRV<sup>S</sup>KDDW PEREMTNS<sup>S</sup>NHLEDPHYSE LTNLKVCIEL TGLHPKKQRH LLHLRERWEQ QVSAADGKPG RQSRKE  
 VTQA TQPEAIPQGT NITEEKGRK RAEAKGN<sup>R</sup>SW SEESLKP<sup>S</sup>SDN EQGLPVFSG<sup>S</sup>PPMKSLS<sup>S</sup>TSS AGGKKQAQPS CAPASRPPAK QQKIKENQ<sup>K</sup>T DVLCAD<sup>E</sup>EED CQAASLL  
 QKY TDNSEKPSGK RLCKTKHLIP QESRRGLPLT GEYVENADG KVTVRRFRKR PEPS<sup>S</sup>DYDLS<sup>S</sup>PAKQEPKPF<sup>D</sup>RLQQLLPASQ STQLPCSSSP QETTQSRPMP PEARRLIVN<sup>K</sup>  
 NAGETLLQRA ARLGYEEVVL YCLENKICDV NHRDNAGYCA LHEACARGWL NIVRHLL<sup>E</sup>YG ADVNCSAQDG TRPLHDAVEN DHLEIVRLL<sup>L</sup>SYGADPTL<sup>A</sup>YSGRTIMKMT HS  
 ELMEKFLT DYLN<sup>D</sup>LQGRN DDDASGTWDF YGSSVCEPDD ESGYDVLANP PGPEDQDDDD DAYSDFE<sup>F</sup>E FSETPLLPCY NIQVSVAQGP RNWLLSDVL KKLKMS<sup>S</sup>RIF RCN  
 FPNVEIV TIAEAEFYRQ VSASLLFSCS KDLEAFNPES KELLDLVEFT NEIQTL<sup>L</sup>GSS VEWLHPSDLA SDNY

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