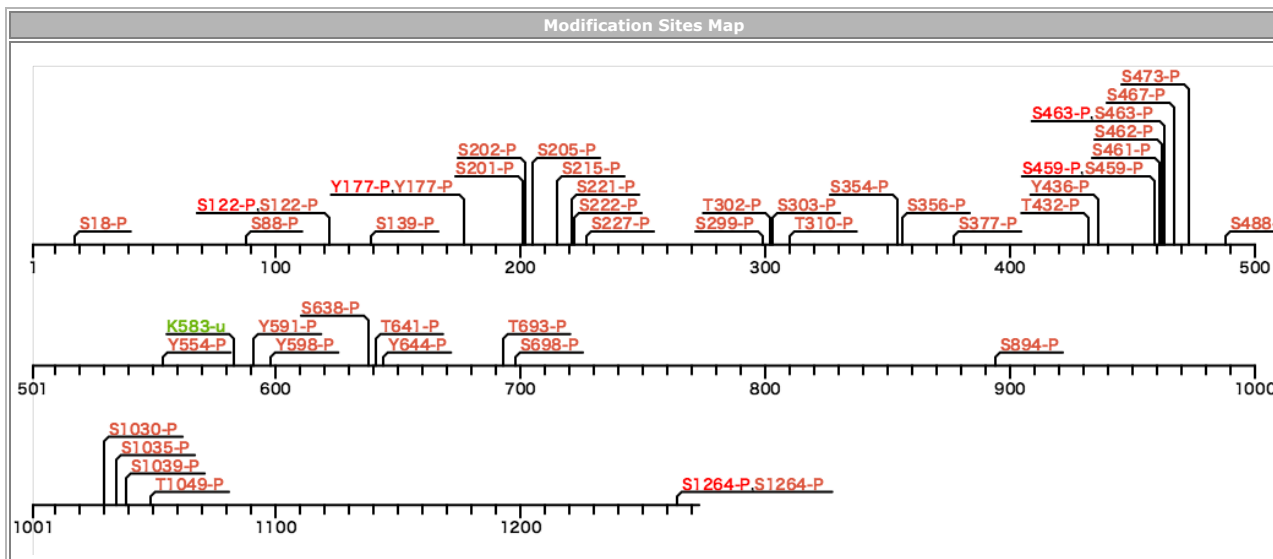


ID	Accession	GeneName	Chr.No.	Description
BCR_HUMAN	P11274	BCR	22q11.23 23521891..23660224	Breakpoint cluster region protein



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
1264	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
1264	S	P	Lab	100627_akimura_pOVICE_2.mgf[F017440]
1264	S	P	Lab	100627_akimura_pOVICE_3.mgf[F017443]
1264	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
1264	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
1264	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
1264	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
1264	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MVDPVGF^uFAEA WKAQFPD^uSEP PRMELRSVGD IEQELERCKA SIRRLEQEVN QERFRMIYLQ TLLAKEKKS^uY DRQRWGFRRRA AQAPDGA^uSEP RASASRPQPA PADGADPPPA E EPEARPDGE G^uSPGKARPGT ARRPGAAA^uSG ERDRGPPAS VAALRSNFER IRKGHGQPGA DAEKPF^uYVNV EFHHERGLVK VNDKEVSDRI S^uLS^uQAMQM ERKKS^uQHGAG S^uSVGDAS^uRPP YRGRSS^uESSC GVDGDYEDAE LNPRFLKDNL IDANGGSRPP WPPLEYQPYQ SIYVGGMMEG EGKGPLLR^uSQ S^uTS^uEQEKRLT^u WPRRSYSPRS FEDCGGGYTP DCSSNENLTS SEEDFSSGQS SRVSP^uPTTY RMFRDKRSP SQNSQS^uFDS SSPPTQCHK RHRHCPVVS EATIVGVRKT GQIWPNDGEG AFHGDADGSF G^uTPPGY^uGCAA DRAEEQRRHQ DGLPYIDDS^uP SSSPHLS^uSKG RGS^uRDALVSG ALESTKA^uSEL DLEKGLEMRK WVLSGILASE ETYLSHLEAL LLPMKPLKAA ATTSQPVLTS QQIETIFFKV PELY^u EIHKEF YDGLFPRVQQ WSHQQRVGD^uL FQK^uLASQLGV Y^uRAFDVNY^uGV AMEMAEKCCQ ANAQFAEISE NLRARSNKDA KDPTTKNS^uLE T^uILLY^uKPVDRV TRSTLVLHDL LKHTP ASHPD HPLLQDALRI SQNFLSSINE EIT^uPRRQS^uSMT VKKGEHRQLL KDSFMVELVE GARKLRHVFL FTDLLCTKL KKQSGGKTQQ YDCKWYIPLT DLSFQMVDEL EAVPNIPL VP DEELDALKIK ISQIKNDIQR EK^uRANKGSKA TERLKKLSE QESLLLLMSP SMAFRVHSRN GKS^uYTFLISS DYERAEWREN IREQKKCFR SFS^uSLTSVELQ MLTNSCVKLQ T VH^uS^uIPLTIN KEDDESPGLY GFLNVIVHSA TGFKQSSNLY CTLEVDSFGY FVNKAKTRVY RDTAEPN^uWNE EFEIELEGSQ TLRILCYEKC YN^uTKIPKED GESTDRLMGK GQVQL DPQAL QDRDWQRTVI AMNGIEVKLS^u VKFNS^uREFSL KRMP^uSRKQT^uG VFGVKIAVVT KRERSKVPYI VRQ^uCV^uEIEIR RGMEEVGIYR VSGVATDIQA LKAAFDVNNK DVSVM SEMD VNAIAGTLKL YFRELPEPLF TDEFYPNFAE GIALSDPVAK ESCMNLNLLS LPEANLLTFL FLLDHLK^uRVA EKEAVNKM^uSL HNLATVFGPT LLR^uSPSEKESK LPANPSQPIT MT DSW^uSLEVM SQVQVLLYFL QLEAIPADS KRQ^uSILFSTE

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