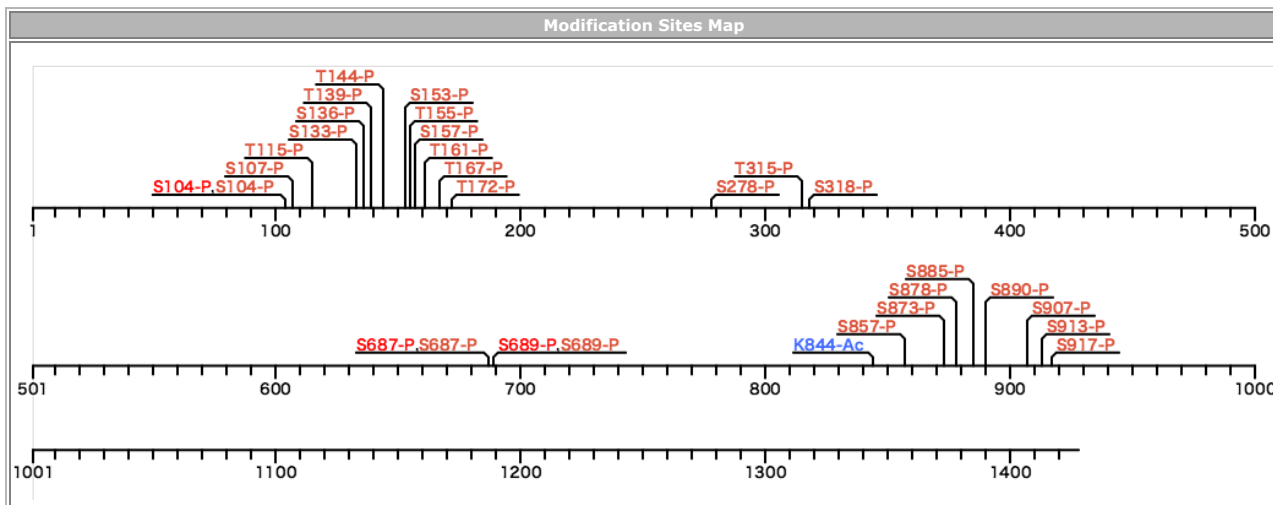


ID	Accession	GeneName	Chr.No.		Description
BCL9_HUMAN	O00512	BCL9	1q21.2	147013182..147098017	B-cell CLL/lymphoma 9 protein



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
687	S	P	Lab	110711_titania_LNCaP_AI_5.mgf[F017446]
687	S	P	Lab	110711_titania_LNCaP_2.mgf[F017434]
687	S	P	Lab	110711_titania_LNCaP_3.mgf[F017435]
687	S	P	Lab	110711_titania_LNCaP_AI_1.mgf[F017441]
687	S	P	Paper	J Proteome Res 2013, 12(1), 260-271

Protein Sequence

MHSSNPKVRS SPNGTQSSP KSKQEVMPVR PTVMSPSGNP QLDSKFSNQG KQGGASQSQ PSPCDKSGG HTPKALPGPG GSMGLKNGAG NGAKGKGKRE RSIADSF
DQ RDPGT PND DS DIKECNSADH IKSQDSQH T P HSM T P SNATA PRSST P SHGQ T TATE T PAQ K T PAKVVVYF STEMANKAAE AVLKGQVETI VSFHIQNISN NKTERSTAPL
NTQISALRND PKPLPQQPPA PANQDQNSSQ NTRLQPTPI PAPAPKPAAP PRPLDRES PG VENKLIPSVG SPASSTPLPP DGTGNSTPN NRAV T P V S QG SNSSSADPKA PPP
PPVSSGE PPTLGENPDG LSQEQLHRE RSLQLRDIQ RMLFPDEKEF TGAQSGGPQG NPGVLDGPQK KPEGPIQAMM AQSQSLGKGP GPRTDVGFAPF GPQGHDRVPF SPD
EMVPPSM NSQSGTIGPD HLDHMTPEQI AWLKLQEFY EEKRRKQEQV VVQCSLQDM MVHQHGPRGV VRGPPPPYQM TPSEGWAPGG TEPFSDGINM PHSLPFRGMA PH
PNMPPGSM RLPGFAGMIN SEMEGPNVPN PASRPGLSGV SWPDDVPKIP DGRNFPPGQG IFSGPGRGER FPNPQGLSEE MFQQQLAEKQ LGLPPGMAME GIRPSMEMNR MI
PGSQRHME PGNNPIFPRI PVEGPL SPSR GDFPKGIPPQ MGPGRLEFG MVPSGMKGDV NLNVNMGNS QMIPQKMREA GAGPEMLKL RPPGSDMLPA QQKMPVLPFG EH
PQQEYGMG PRPFLPMSQG PGNSGLRNL REPIGPDQRT NSRSLHMPPL PLNPSSNPTS LNTAPPVQRG LGR K PLDISV AGSQVH S PGI NPLKSPTMHQ VQ S PMLG S PS GNL
K S P Q T P S QLAGMLAGPA AAASIK SPPV LG S AAA S P V H LKSPSLPAPS PGWTSSPKPP LQSPGIPPNH KAPLTMASPA MLGNVESGGP PPPTASQPAS VNIPGSLPSS TPYTMP
PEPT LSQNPLSIMM SRMSKFAMPS STPLYHDAIK TVASSDDDSP PARSPNLPSM NNMPGMGINT QNPRISGPNP VVPMPTLSPM GMTQPLSHSN QMPSNAVGP NIPPHGV
PMG PGLMHNPIM GHGSQEPMPV PQRGMGFPQG FPPVQSPQQ VPFPHPNGPSG GQGSFPGGMG FPEGGLGRP SNLPQSSADA ALCKPGGPGG PDSFTVLGNS MPVSFT
DPDL QEVIRPGATG IPEFDLSRII PSEKPSQTLQ YFPRGEVPGR KQPQGGPGF SHMQGMMGEQ APRMGLALPG MGGPGVGTPT DIPLGTAPSM PGHNPMRPPA FLQQGMM
GPH HRMMSPAQST MPGQPTLSMN PAAAVGMIPG KDRGPAGLYT HPGVPGSPGM MMSMQGMMGP QQNIMIPQM RPRGMAADVG MGGFSQGGPN PGNMM

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