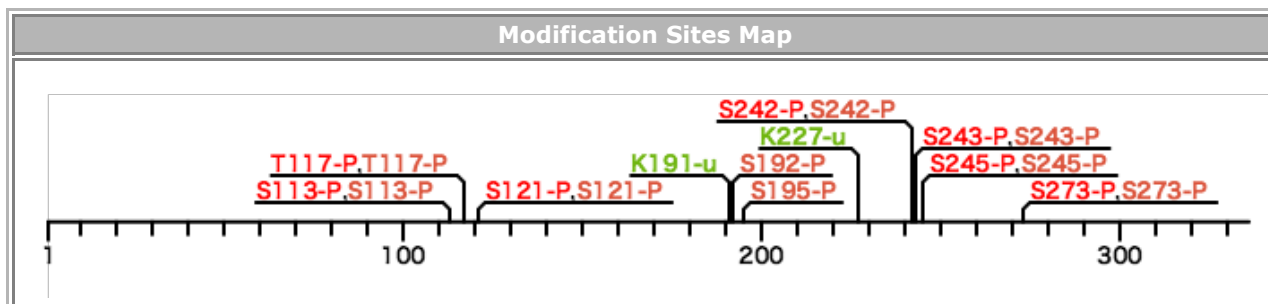


ID	Accession	GeneName	Chr.No.		Description
B2L12_HUMAN	Q9HB09	BCL2L12	19q13.33	50168823..50177173	Bcl-2-like protein 12



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
242	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
242	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
242	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
242	S	P	Lab	140320_Agarose_.mgf[F017423]
242	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
242	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
242	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
242	S	P	Lab	110218_pRMG2_1.mgf[F017475]
242	S	P	Lab	110218_pRMUGS_1.mgf[F017479]
242	S	P	Lab	100627_akimura_pOVISE_1.mgf[F017437]
242	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
242	S	P	Lab	100627_akimura_pOVTOKO_1.mgf[F017447]
242	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
242	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
242	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
242	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
242	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
242	S	P	Paper	Mol Cell Proteomics 2010, 9(7), 1540-1553
242	S	P	Paper	Proc Natl Acad Sci USA 2008, 105(38), 10762-10767
242	S	P	Paper	Sci Signal 2010, 3(104), ra3
242	S	P	Paper	Sci Signal 2011, 4(164), rs3

**Protein Sequence**

MGRPAGLFPP LCPFLGFRPE ACWERHMQIE RAPSVPFLR WAGYRPGPVR RRGKVELIKF VRVQWRRPQV EWRRR  
 RWGPG PGASMAGSEE LGLREDTLRV LAAFLRRGEA AGSPVPTPPR SPAQEPTDF LSRLRRLPC SLGRGAAPSE  
 SPRPCSLPIR PCYGLEPGA TPDFYALVAQ RLEQLVQEQL KSPPSPELQG PPSTEKEAIL RRLVALLEE AEVINQKLA  
 S DPALRSKLVR LSSDSFARLV ELFCSRDDSS RPSRACGPP PPSPEPLARL ALAMELSRRV AGLGGTLAGL SVEHV  
 HSFTP WIQAHGGWEG ILAVSPVDLN LPL

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