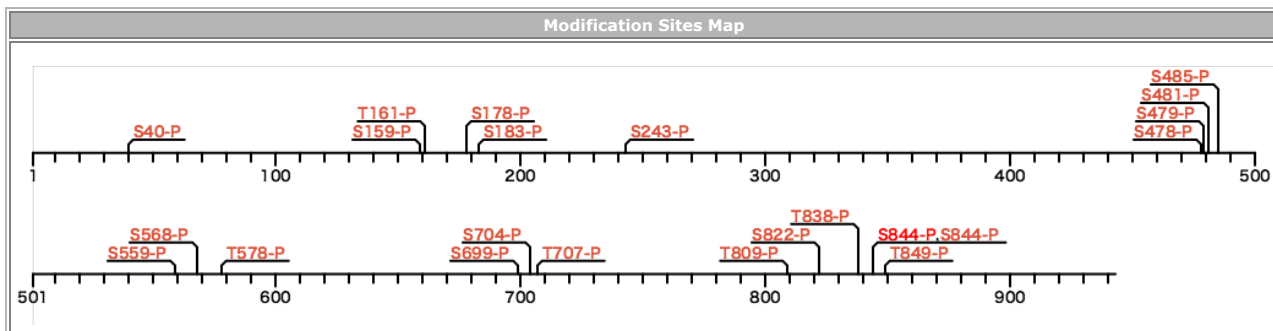


ID	Accession	GeneName	Chr.No.		Description
BBX_HUMAN	Q8WY36	BBX	3q13.12	107241783..107530171	HMG box transcription factor BBX



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
844	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
844	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
844	S	P	Lab	140320_Agarose_.mgf[F017423]
844	S	P	Lab	110218_pRMG2_1.mgf[F017475]
844	S	P	Lab	110218_pRMG2_3.mgf[F017477]
844	S	P	Lab	110218_pRMG2_4.mgf[F017478]
844	S	P	Lab	110711_titania_LNcaP_AI_3.mgf[F017444]
844	S	P	Lab	110711_titania_LNcaP_1.mgf[F017433]
844	S	P	Lab	110711_titania_LNcaP_4.mgf[F017436]
844	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
844	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
844	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
844	S	P	Lab	100520-GIST-R2.mgf[F017517]
844	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
844	S	P	Lab	100520-GIST-R3.mgf[F017519]
844	S	P	Lab	100520-GIST-W2.mgf[F017522]
844	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
844	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
844	S	P	Paper	J Proteomics 2014, 96, 253-262
844	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MKGSNRNKDH SAEGEGVGKR PKRKCLQWHP LLAKLLDFSEEEEEDEEE DIDKVQLLGA DGLEQDVGET EDDESPEQRA RRPNNAFLF CKHRSLVRQ EHPRLDNRGA TKILADWWAV LDPKEQKYT DMAKEYKDAF MKANPGYKWC PTTNPKVKSPTPTVNP RKKL WAFPSDS SRD LPSPKKAKTE EMPQLNFGMA DPTQMGGLSM LLLAGEHALG TPEVSSGTCR PDVSEPELR QKSP LFQFAE ISSSTSHSDA STKQCQTSAL FQFAEISSNT SQLGGAEPVK RCGKSALFQL AEMCLASEGM KMEESKLIKA KESDGGRIKE LEK GKEEKI KMEKTDETRL QKEAEFEKSA KENLRDSKEL RNFEALQIDD IMAIKMEDPK EIRKEELEED HKCSHFDFE YSASSKIIS DVPSRKDHMC HPHGIMIIED PAALNKPE KL KKKKKSKMD RHGNDKSTPK KTCKKRQSSDIE SVIYTI EAVAKGDWGI EKLGDTPRKK VRTSSSGKGS ILDAKPPKKK VKSREKKMSK EKSSD TTKE RPPDFIS SA SKNISGETPE GIKAEPLTPM EDALPPSLSG QAKPEDSDCH RKIETCGSRK SERSCKGALY KTLVSEGMLT SLRANVDRGK RSSGKGNSSD HEGCWNEESW TFSQSGTSGS KKFKKTPKPE DCLLGS AKLD EEFKFKFNSL PQYSPVTFDR KCVPVPRKKK KTGNSVSEPT KTSKGFQSQ KKNLFHKIVS KYKHKKEKPN VPEKSGGDKW SNKQLFLDAI HP TEAIFSED RNTMEPVHKV KNIPSIFNTP EPTTTQEP LVGSQKRKARKT KITHLVRTAD GRVSPAGGTL DDKPKQLQR SLPKATETDC NDKCSHNTVE GETRSSTPEM PAVSA FFSLA ALAEVAAMEN VHRGQRSTPL THDGQPKEMP QAPVLISCAD

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