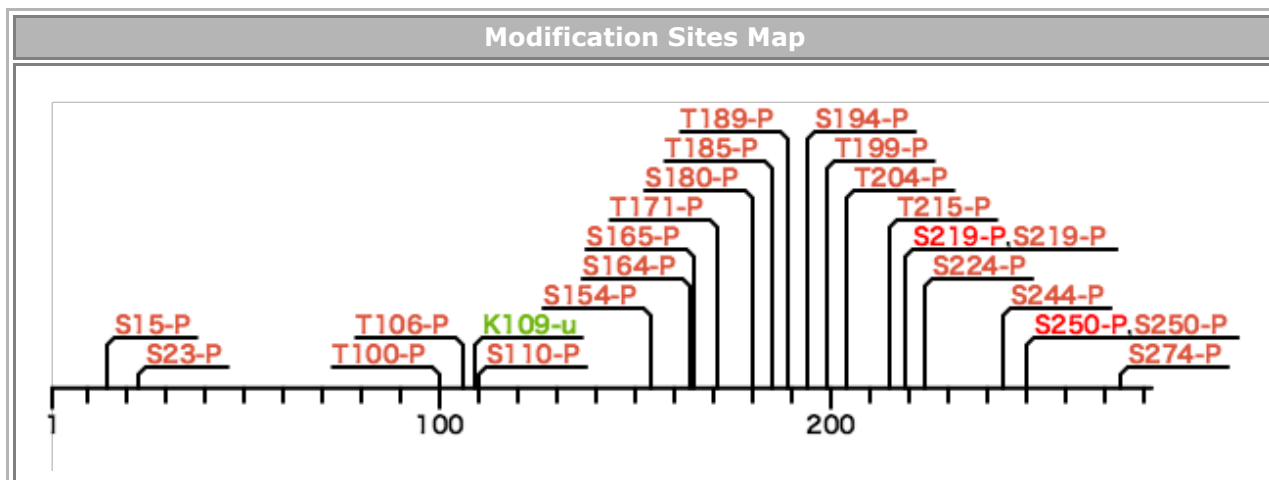


ID	Accession	GeneName	Chr.No.		Description
BOREA_HUMAN	Q53HL2	CDCA8	1p34.3	38158090..38175391	Borealin



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
219	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
219	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
219	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
219	S	P	Lab	140320_Agarose_.mgf[F017423]
219	S	P	Lab	140320_Agarose_.mgf[F017423]
219	S	P	Lab	110218_pRMUGS_2.mgf[F017480]
219	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
219	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
219	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
219	S	P	Lab	100628_akimura_pOVSAHO_2.mgf[F017461]
219	S	P	Lab	100628_akimura_pOVSAHO_3.mgf[F017462]
219	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
219	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
219	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
219	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
219	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
219	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
219	S	P	Lab	110218_pOVSAYO_1.mgf[F017469]
219	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
219	S	P	Lab	110218_pRMG1_1.mgf[F017472]
219	S	P	Lab	110218_pRMG1_3.mgf[F017474]
219	S	P	Lab	110218_pRMG2_1.mgf[F017475]

219	S	P	Lab	110218_pRMG2_2.mgf[F017476]
219	S	P	Lab	110218_pRMG2_3.mgf[F017477]
219	S	P	Lab	110218_pRMG2_4.mgf[F017478]
219	S	P	Lab	110218_pRMUGS_1.mgf[F017479]
219	S	P	Lab	100627_akimura_pOVISe_1.mgf[F017437]
219	S	P	Lab	100627_akimura_pOVISe_2.mgf[F017440]
219	S	P	Lab	110711_titania_LNcaP_AI_2.mgf[F017442]
219	S	P	Lab	100627_akimura_pOVISe_3.mgf[F017443]
219	S	P	Lab	110711_titania_LNcaP_AI_3.mgf[F017444]
219	S	P	Lab	110711_titania_LNcaP_AI_4.mgf[F017445]
219	S	P	Lab	110711_titania_LNcaP_AI_5.mgf[F017446]
219	S	P	Lab	110711_titania_LNcaP_AI_6.mgf[F017448]
219	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
219	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
219	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
219	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
219	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
219	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
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219	S	P	Lab	140320_tita_C18_.mgf[F017426]
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219	S	P	Lab	110711_titania_LNcaP_AI_1.mgf[F017441]
219	S	P	Lab	100520-GIST-IM1.mgf[F017509]
219	S	P	Lab	100520-GIST-IM2.mgf[F017512]
219	S	P	Lab	100520-GIST-IM3.mgf[F017514]
219	S	P	Lab	100520-GIST-R2.mgf[F017517]
219	S	P	Lab	100520-GIST-R3.mgf[F017519]
219	S	P	Lab	100520-GIST-W1.mgf[F017521]
219	S	P	Lab	100520-GIST-W2.mgf[F017522]
219	S	P	Lab	100520-GIST-W3.mgf[F017524]
219	S	P	Paper	Cancer Res 2009, 69(6), 2663-2668
219	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594

219	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
219	S	P	Paper	Mol Cell 2008, 31(3), 438-448
219	S	P	Paper	Mol Cell Proteomics 2009, 8(7), 1751-1764
219	S	P	Paper	Proc Natl Acad Sci USA 2008, 105(38), 10762-10767
219	S	P	Paper	Sci Signal 2009, 2(84), ra46
219	S	P	Paper	Sci Signal 2010, 3(104), ra3

Protein Sequence

MAPRKGSSRV AKTNSLRRRK LASFLKDFDR EVEIRIKQIE SDRQNLLKEV DNLYNIEILR LPKALREMNW
LDYFALGGNK QALEEAATAD LDITEINKLT AEAIQTPLKS AKTRKVIQVD EMIVEEEEEEE ENERKNLQTA R
VKRCPPSKK RTQSIQGKGGK GKRSSRANTV TPAVGRLEVS MVKPTPGLTP RFD~~S~~RVFK~~T~~P GLR~~T~~PAAGER
IYNI~~S~~GNG~~S~~P LAD~~S~~SKEIFLT VPVGGGESLR LLASDLQRHS IAQLDPEALG NIKKLSNRLA QIC~~S~~SIRTH

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