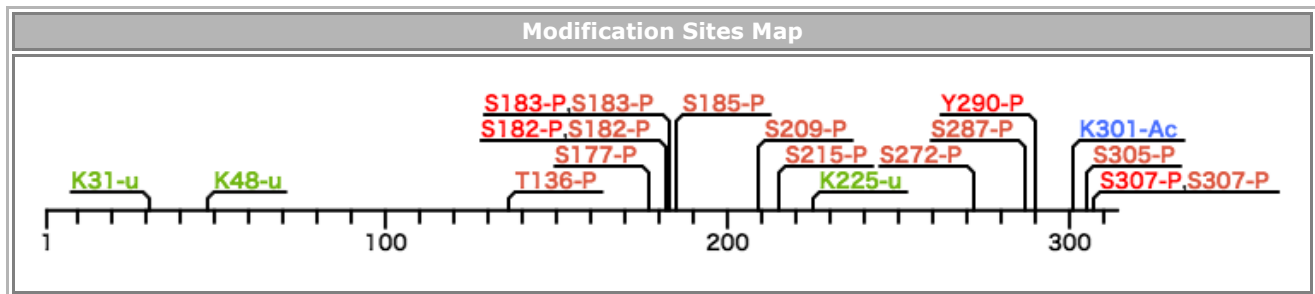


ID	Accession	GeneName	Chr.No.		Description
CPIN1_HUMAN	Q6FI81	CIAPIN1	16q21	57462081..57481440	Anamorsin



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
183	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
183	S	P	Lab	110218_pRMG2_3.mgf[F017477]
183	S	P	Lab	100627_akimura_pOVISE_1.mgf[F017437]
183	S	P	Lab	110711_titania_LNCaP_AI_4.mgf[F017445]
183	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
183	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
183	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
183	S	P	Lab	110711_titania_LNCaP_2.mgf[F017434]
183	S	P	Lab	110711_titania_LNCaP_3.mgf[F017435]
183	S	P	Lab	110711_titania_LNCaP_5.mgf[F017438]
183	S	P	Lab	100520-GIST-IM1.mgf[F017509]
183	S	P	Lab	100520-GIST-IM3.mgf[F017514]
183	S	P	Lab	100520-GIST-R3.mgf[F017519]
183	S	P	Paper	Cancer Res 2009, 69(6), 2663-2668
183	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
183	S	P	Paper	J Proteome Res 2009, 8(11), 4954-4965
183	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
183	S	P	Paper	Mol Cell Proteomics 2011,10(11), M111.008540
183	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MADFGISAGQ FVAVVWDKSS PVEALKGLVD **K**LQALGTNEG RVSVENI**K**QL LQSAHKESSEF DIILSGLVPG STTLHSAEIL AEI
 ARILRPG GCLFLKEPVE TAVDNNSKVK TASKLCSALT LSGLVEVKEL QREPL**T**PEEV QSVREHLGHE SDNLLFVQIT GKPPNF
 EVGS SRQLK**L**SITK **KSS**PSVKPAV DPAAAKLWTL SANDMEDD**S**M DLID**S**DELLD PEDL**K**KPDPA SLRAASCCEG KKRKACK
 NCT CGLAELEKE KSREQMSSQP **K**SACGNCYLG DAFRCAS**C**PY LGMPAFKPGE **KVLL**S**D**S**N**LH DA

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