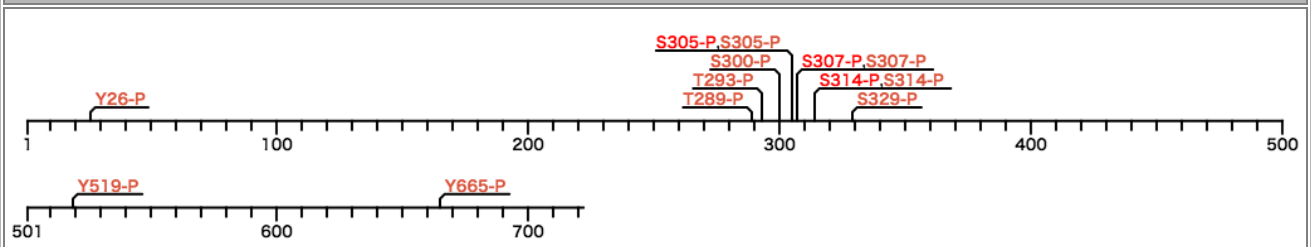
ID	Accession	GeneName	Chr.No.		Description
CNKR1_HUMAN	Q969H4	CNKSR1	1p36.11	26503894..26516377	Connector enhancer of kinase suppressor of ras 1

Modification Sites Map



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
305	S	P	Lab	140320_Agarose_.mgf[F017423]
305	S	P	Lab	110218_pRMUGS_2.mgf[F017480]
305	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
305	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
305	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
305	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
305	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
305	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
305	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
305	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
305	S	P	Lab	110218_pOVSAYO_1.mgf[F017469]
305	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
305	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
305	S	P	Lab	110218_pRMG1_1.mgf[F017472]
305	S	P	Lab	110218_pRMG1_2.mgf[F017473]
305	S	P	Lab	110218_pRMG2_1.mgf[F017475]
305	S	P	Lab	110218_pRMG2_3.mgf[F017477]
305	S	P	Lab	110218_pRMG2_4.mgf[F017478]
305	S	P	Lab	100627_akimura_pOVISE_1.mgf[F017437]
305	S	P	Lab	100627_akimura_pOVISE_2.mgf[F017440]
305	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
305	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
305	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
305	S	P	Lab	100627_akimura_pRMG1_3.mgf[F017453]
305	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
305	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
305	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
305	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
305	S	P	Lab	140320_tita_C18_.mgf[F017426]
305	S	P	Lab	140320_tita_SDB_.mgf[F017430]
305	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
305	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
305	S	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668
305	S	P	Paper	Mol Cell Proteomics 2014, 13(7), 1690-1704

Protein Sequence

MEPVETWTPG KVATWLRGLD DSLQDYPFED WQLPGKNLLQ LCPQSLEALA VRSLGHQELI LGGVEQLQAL SSRLQTENLQ SLTEGLLGAT HDFQSIVQGC LGDCAKTPID VL CAAVELLH EADALLFWLS RYLFSHLNDF SACQEIRDLL EELSQVLHED GPAAEKEGTV LRICSHVAGI CHNILVCCPK ELLEQKAVLE QVQLDSPLGL EIHTTSNCQH FVSQVD TQVP TDSRLQIQPG DEVVQINEQV VVREERDMVG WPRKNMVREL LREPAGLSLV LKKIPIPETP PQTPPQVLDS PHQRSPSLSL APLSPRAPSE DVFAFDLSSN PSPGPSPAW T DSASLGPEPL PIPPEPPAIL PAGVAGTPGL PESPDKSPVG RKKSKGLATR LSRRRVSCRE LGRPDCDGWL LLRKAPGGFM GPRWRRRWFV LKGHTLYWYR QPQDEKAEGL INVSNYSLES GHDQKKKYVF QLTHDVYKPF IFAADTLTDL SMWVRHLITC ISKYQSPGRA PPPREEDCYS ETEAEDPDDE AGSHSASPSP AQAGSPLHGD TSPAATPTQR SP RTSFGSLT DSSEEALEGM VRGLRQGGVS LLGQPQPLTQ EQWRSSFMRR NRDPQLNERV HRVRALQSTL KAKLQELQVL EEVLGDPELT GEKFRQWKEQ NRELYSEGLG AW GVAQAEGS SHILTSDSTE QSPHSLPSDP EEHSHLCPLT SESSLRPPDL

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