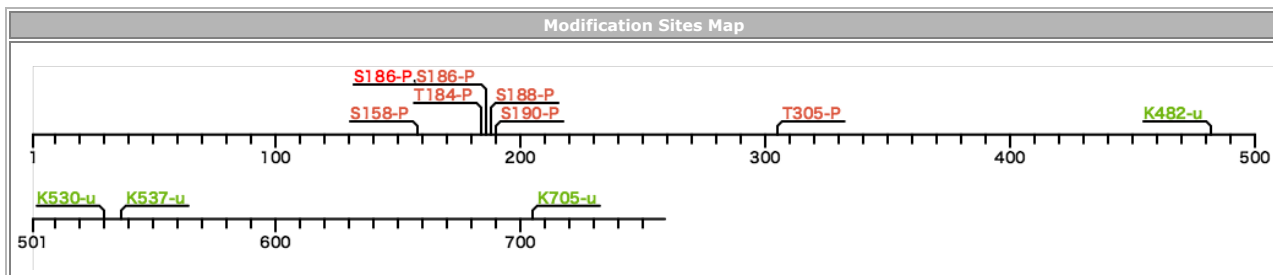


ID	Accession	GeneName	Chr.No.	Description
CCPG1_HUMAN	Q9ULG6	CCPG1	15q21.3 55632230..55700708	Cell cycle progression protein 1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
186	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
186	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
186	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
186	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
186	S	P	Paper	J Proteomics 2014, 96, 253-262
186	S	P	Paper	Mol Cell Proteomics 2015, 14(6), 1599-1615
186	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MSENSSDSDS SCGWTVISHE GSDIEMLSNV TPTDSECPAP ECSSLEQEEL QALQIEQGES SQNGTVLMEE TAYPALEETS STIEAEEQKI PEDSIYIGTA SDDSDIVTLE PPKL EEIGNQ EVVIVIEEAQS SEDFNMGSSS SSQYTFQCPE TVFSSQP^SDD ESSSDETS^SNQ PSPAFRRRRR RKK^TV^SA^SE^S EDRLVAEQET EPSKELSKRQ FSSGLNKCVI LALVIAI SMG FGHFYGTIQI QKRQQLVRKI HEDELNDMKD YLSQCQQEQE SFIDYKSLKE NLARCVLTLTE AEKMSFETQK TNLAT^TENQYL RVSLEKEEKA LSSLQEELNK LREQIRILED KGTSTELVKE NQKLKQHLEE EKQKKHSFLS QRETLLEAK MLKRELERER LVTTLALRGEL QQLSGSQLHG KSDSPNVYTE KKEIAILRER LTELERKLT^F EQQRSDLWER LYVE AKDQNG KQGTGDKKKK GRGSHRAK^NSK^EETFLGSVK ETFDAMKNST KEFVRHHKEK IKQAKEAVKE NLKKFSDSV^K STFRHF^KDTT KNIFDEKGNK RFGATKEAAE KPRT VFSDYL HPQYKAPTEN HHNRGPTMQN DGRKEKPVHF KEFRKNTNSK KCSPGHDCRE NSHSFRKACS GVFDCAQ^SES MSLFNTVVNP IRMDEF^RQII QRYMLKELDT FCHW NELDQF INKFLNGVF IHDQKLFTDF VNDV^KDYLRN MKEYEVDNDG VFEKLDEYIY RHFFGHTFSP PYGPRSVYIK PCHYSS

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