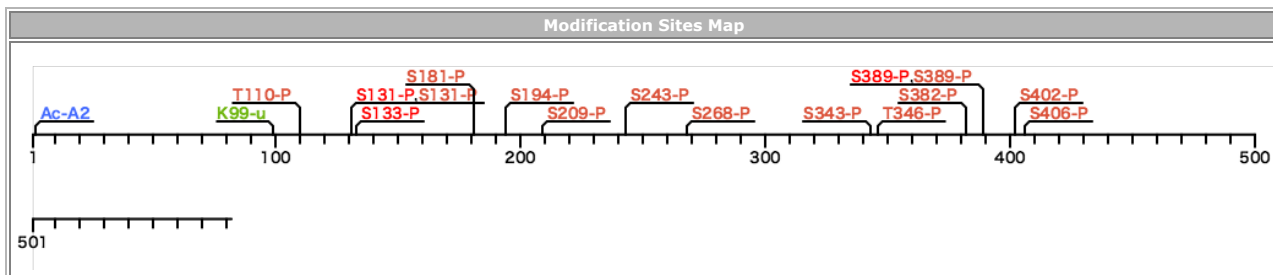


ID	Accession	GeneName	Chr.No.		Description
CARF_HUMAN	Q9NXV6	CDKN2AIP	4q35.1	184365744..184369351	CDKN2A-interacting protein



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
131	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
131	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
131	S	P	Lab	140320_tita_C18_.mgf[F017426]
131	S	P	Lab	140320_tita_SDB_.mgf[F017430]
131	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
131	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
131	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
131	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
131	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
131	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
131	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
131	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
131	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MAQEVSEYLS QNPRVAAWVE ALRCDGETDK HWRHRRDFLL RNAGDLAPAG GAASASTDEA ADAESGTRNR QLQQLISFSM AWANHVFLLGC RYPQKVMDKI LSMAEGIKVT
 DAPTYTTRDE LVAKVKKRGI SSSNEGVEEP SKKRVIIEGKN SSAVEQDHAK TSAKTERASA QQENSSTCIG SAIKSESGNS ARSSGISSQN SSTSDGDRSV SSQSSSSVSS Q
 VTTAGSGKA SEAEAPDKHG SASFVSLKLS SVNSHMTQST DSRQQSGSPK KSALEGSSAS ASQSSSEIEV PLLGSSGSSE VELPLLSSKP SSETASSGLT SKTSSEASVS SSV
 AKNSSSS GTSLTPKSS SSTNTSLLTS KSTSQAASL LASKSSSQTS GSLVSKTSL ASVSQLASKS SSQTSTSQLP SKTSQSSES SVKFSCKLTN EDVKQKQPPF NRLYKT
 VAWK LVAVGGFSPN VNHGELLNAA IEALKATLDV FVPLKELAD LPQNKSSQES IVCELRCVSK YLGTGCGKSK ENAKAVASRE ALKFLKLVK VVKICKRKYR GSEIEDLVLL
 DEESRPVNL PALKHPQEL

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