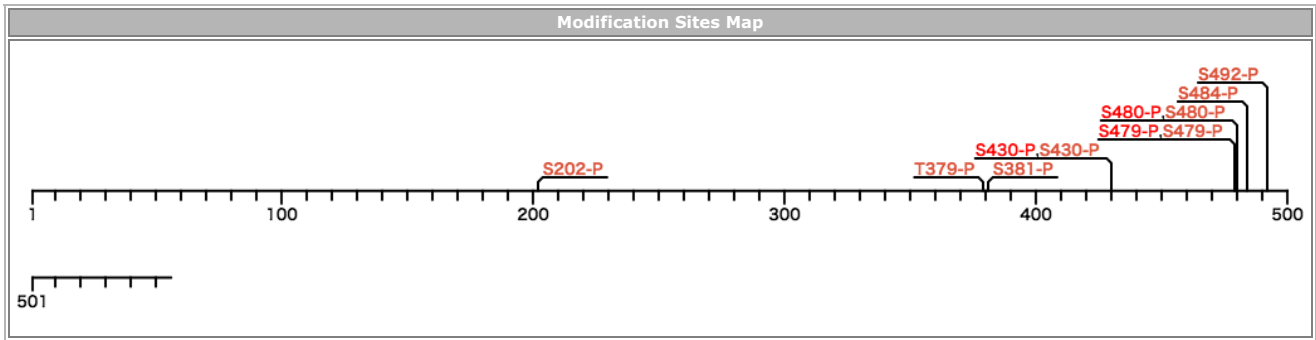


ID	Accession	GeneName	Chr.No.	Description
DNJC1_HUMAN	Q96KC8	DNAJC1	10p12.31 22045466..22292698	DnaJ homolog subfamily C member 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
430	S	P	Lab	100520-GIST-IM1.mgf[F017509]
430	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
430	S	P	Lab	100520-GIST-IM2.mgf[F017512]
430	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
430	S	P	Lab	100520-GIST-IM3.mgf[F017514]
430	S	P	Lab	100520-GIST-R2.mgf[F017517]
430	S	P	Lab	100520-GIST-R3.mgf[F017519]
430	S	P	Lab	100520-GIST-W2.mgf[F017522]
430	S	P	Lab	100520-GIST-W3.mgf[F017524]
430	S	P	Paper	Sci Signal 2009, 2(84), ra46
430	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MTAPCSQPAQ LPGRRQLGLV PFP PPPPRTP LLWLLLLLLA AVAPARGWES GDLELFDLVE EVQLNFYQFL GVQQDASSAD IRKAYRKLSL TLHPDKNKDE NAETQFRQLV AIYE VLKDDE RRQRYDDILI NGLPDWRQPV FYRRVRKMS NAELALLFI ILTVGHYAVV WSIYLEKQLD ELLSRKKREK KKKTGSKSVD VSKLGASEKN ERLLMKPQWH DLLPCKL GIW FCLTLKALPH LIQDAGQFYA KYKETRLKEK EDALTRTELE TLQKQKVKK PKPEFPVYTP LETTYIQSYD HGTSIEEIEE QMDDWLENRN RTQKKQAPEW TEEDLSQLTR SMVKFPGGTP GRWEKIAHEL GRSVTDVTTK AKQLKDSVTCSPGMVRLSEL KSTVQNSRPI KTATLPPDM ITQREDAEGV AAEFEQEGDS GEQETGATDA RPRRRKPARL L EATAKPEPE EKSRAKRQKD FDIAEQNES DEESLRKERA RSAEEPWTQN QKLELALQ QYPRGSSDRW DKIARCVPSK SKEDCIARYK LLVELVQKKK QAKS

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