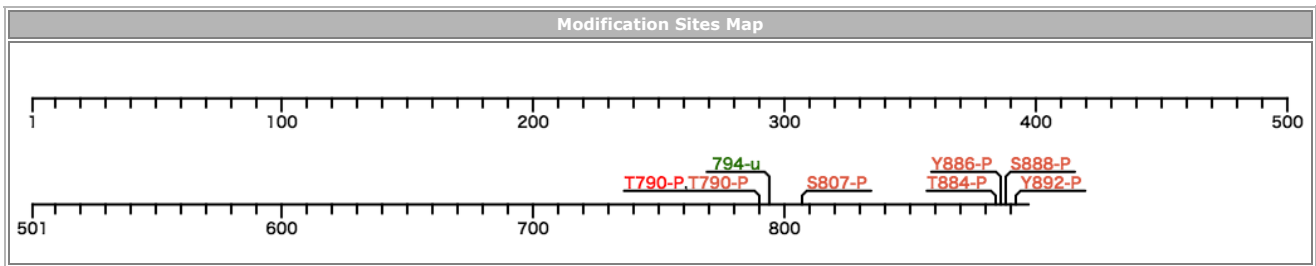


ID	Accession	GeneName	Chr.No.	Description
DAG1_HUMAN	Q14118	DAG1	3p21.31 49506146..49573048	Dystroglycan



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
790	T	P	Lab	130327_HEK_CE_pphos.mgf[F015007]
790	T	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
790	T	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MRMSVGLSLL LPLSGRTFLL LLSVVMAQSH WPSEPSEAVR DWENQLEASM HSVLSDLHEA VPTVVGIPDG TAVVGRSFRV TIPTDLIASS GDIKVSAAK KEALPSWLHW DS QSHTLEGL PLDTDKGVHY ISVSATRLGA NGSHIPQTSS VFSIEVYPED HSELQSVRTA SPDPGEVSS ACAADPEPTV LTVILDADLT KMPKQRIDL LHRMRSFSEV ELHNM KLV PV VNNRFLDMSA FMAGPGNAKK VVENGALLSW KLGCSLNQNS VPDIHGVEAP AREGAMSAQL GYPVVGWHIA NKKPPLPKRV RRQIHATPTP VTAIGPPTTA IQEPPSR IVP TPTSPAIAAPP TETMAPPVRD PVP GKPTVTI RTRGAIITQP TLGPIQPTRV SEAGTTVPGQ IRPTMTIPGY VEPTAVATPP TTTTKKPRVS TPKPATPSTD STTTTTRRPT KKPR TPRPVP RVTTKVSITR LETASPTRI RTTTSVGPVRG GEPNQRPELK NHIDRVDRAWV GTFYEVKIPS DTFYDHEDTT TDKLKLTLKL REQQLVGEKS WVQFNSNSQL MYGLPDS SHV GKHEYFMHAT DKGGLSAVDA FEIHVHRRPQ GDRAPARFKA KFVGDPALVL NDIHKKIALV KKLAFAFGDR NCSTITLQNI TRGSIVVEWT NNTLPLEPCP KEQIAGLSRR IAEDDGKPRP AFSNALEPDF KATSITVTGS GSCRHLQFIP VPPRRVPSE APTEVPDRD PEKSEDDVY LHTVIPAVVV AAILLIAGII AMICYRKKRK GKLTLEDQAT FIKKGV PIIF ADELDDSKPP PSSSMLILQ EEKAPLPPPE YPNQSVPETT PLNQDTMGEY TPLRDEDNA PPYQPPPPPT APMEGKSRP KNMTPYRSPP PYVPP	

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