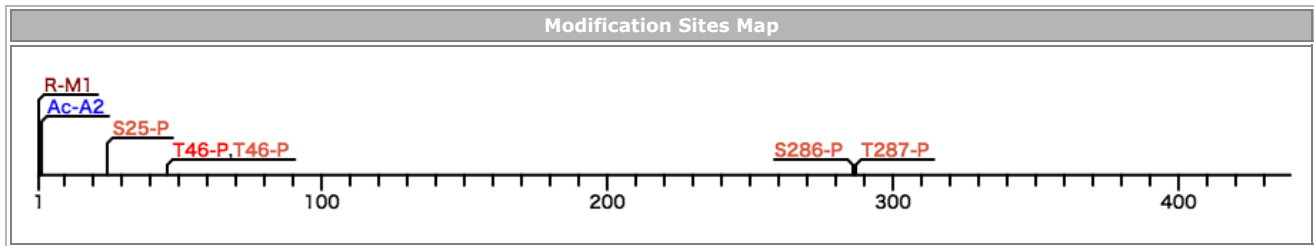


ID	Accession	GeneName	Chr.No.		Description
EF1G_HUMAN	P26641	EEF1G	11q12.3	62327073..62342401	Elongation factor 1-gamma



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
46	T	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
46	T	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MAAGTLYTYP ENWRAFKALI AAQYSGAQVR VLSAPPHFH GQTNRTPEFL RKFPAGKVPA FEGDDGFCVF ESNAIYYVS NEELRGSTPE AAAQVVQW
 VS FADSDIVPPA STWVFPTLGI MHHNKQATEN AKEEVRRILG LLDAYLKTRT FLVGERVTLA DITVCTLLW LYKQVLEPSF RQAFPNTNRW FLTCINQP
 QF RAVLGEVKLC EKMAQFDACK FAETQPKKDT PRKEKGSREE KQKQAERKE EKKAAAPAE EEMDECEQAL AAEPKAKDPF AHLPKSTFVL DEFKRK
 YSNE DTLSVALPYF WEHFDKDGWS LWYSEYRFPE ELTQTFMSCN LITGMFQRLD KLRKNAFASV ILFGTNNSSS ISGVVWFRGQ ELAFPLSPDW QVD
 YESYTW R KLDPGSEETQ TLVREYFSWE GAFQHVKGKAF NQGKIFK

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