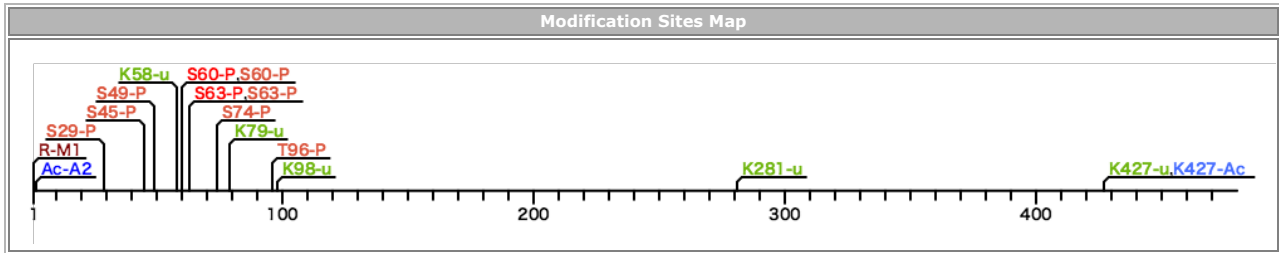


ID	Accession	GeneName	Chr.No.		Description
AMP22_HUMAN	P50579	METAP2	12q22	95867296..95909615	Methionine aminopeptidase 2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
60	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
60	S	P	Paper	Proc Natl Acad Sci USA 2008, 105(38), 10762-10767
60	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
60	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
60	S	P	Paper	Mol Cell Proteomics 2010, 9(7), 1540-1553

Protein Sequence

MA GVVEEVAAS GSHLNGDLDP DDREEGAAS T AEEAAKKRR KKKKSKGPSA AGEQEPDKES GASVDEVARQ LERSALEDKE RDEDDGDG DGDGATGKK KKKKKK
 RGPV QTDPPSVPI CDLYPNGVFP KGQCEYPT QDGRTAAWRT TSEEKALDQ ASEEIWNDFR EAAEAHRQVR KYVMSWIKPG MTMIEICEKL EDCSRKLIKE NGLN
 AGLAFP TGCSLNNCAA HYPNAGDTT VLQYDDICKI DFGTHISGRI IDCAFTVTFN PKYDILLKAV KDATNTGIKC AGIDVRLCDV GEAIQEVMS YEVEIDGKTY QVKPI
 RNLNG HSGIQYRIHA GKTVPVKGKGG EATRMEEGEV YAIETFGSTG KGVVHDDMEC SHYMKNFVVG HVPRLRPTK HLLNVINENF GTLAFCCRRL DRLGESK YLM ALKN
 LCDLGI VDPYPLCDI KGSYTAQFEH TILLRPTCKE VVSRGDD

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