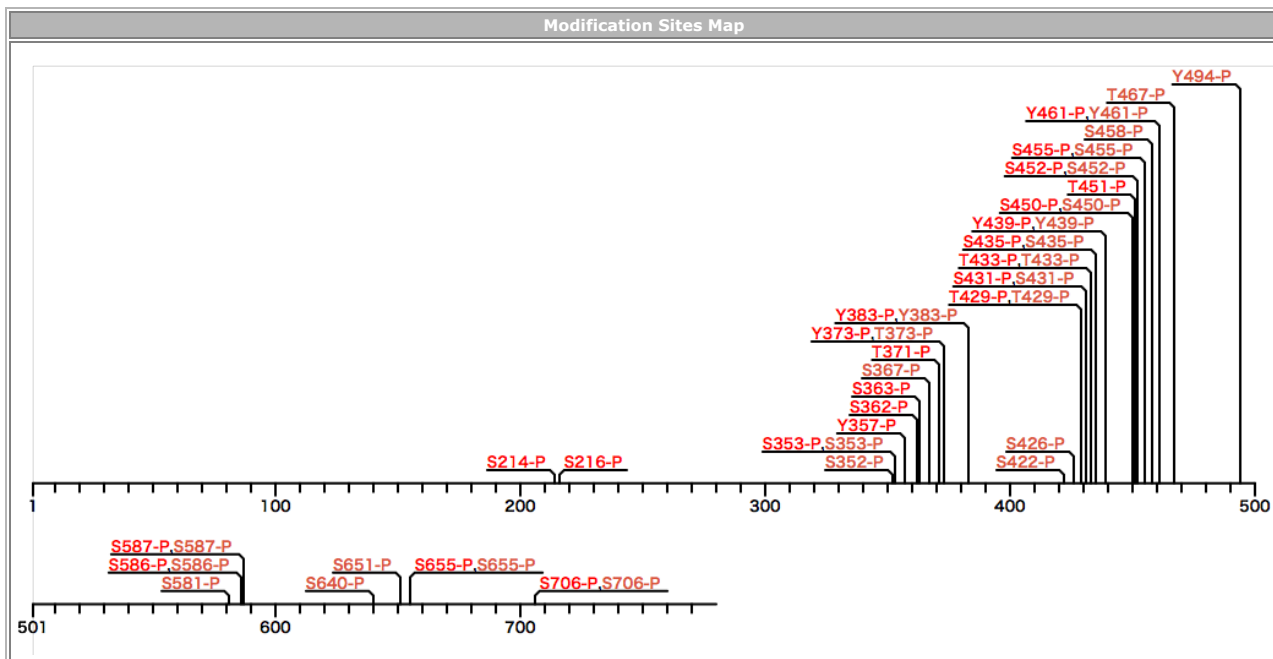


ID	Accession	GeneName	Chr.No.	Description
ABLM1_HUMAN	O14639	ABLM1	10q25.3 116190872..116444762	Actin-binding LIM protein 1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
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Protein Sequence	
MPAFLGLKCL GKLCSSEKSK VTSSERTSAR GSNRKRRLIVE DRRVSGTSFT AHRRTITLH LYLCPKDYCP RGRVCNSVDP FVAHPQDPHH PSEKPVHICH KCGEPCKGEV LRV	QTKHFHI KCFTCKVCGC DLAQGGFFIK NGEYLCTLDY QRMYGTRCHG CGEFVEGEV TALGKTYHPN CFACTICKRP FPPGDRVTFN GRDCLCLQCA QPMS ^{SS} PKET TFSSN
CAGCG RDIKNGQALL ALDKQWHLGC FKCKSCGKVL TGEYISKDGA PYCEKDYQGL FGVKCEACHQ FITGKMLEAG DKHYHPSCAR CSRCNQMFTE GEEMYLQGST VWHP	DCKQST KTEEKLRPTR T ^{SS} ESI ^Y SRP G ^{SS} IPGSPGH T ^{IY} AKVDNEI LD ^Y KDLAIP KVKAIYDIER PDLITYEPFY TSGYDDKQER Q ^{SL} GES ^{SP} RL ^T S ^{PT} PS ^{AE} GYQ D ^{VR} DRMIHR
S ^{TS} QGS ^{IN} SPV YSRHSYTP ^{TT} SRSPQHFRP GNEPSSGRNS PL ^Y RPDSRP LTPTYAQAPK HFHVPDQGIN IYRKPPIYKQ HAALAAQSKS SEDIIKFSKF PAAQAPDPSE TPK	IETDHW P ^{PP} SFAVVG P ^{DM} KRR ^{SS} GRE EDDEELLRRR QLQEEQLMKL NSGLGQLILK EEMEKESRER SLLASRYD ^S PINSASHIPS S ^K TAS ^L PGYG RNLGHRPVST DFAQY
NSYGD VSGGVRDYQT LPDGHMPAMR MDRGV ^S MPNM LEPKIFPYEM LMVTNRGRNK ILREVDTRL ERHLAPEVFR EIFGMSIQEF DRLPLWRRND MKKKAKL	

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