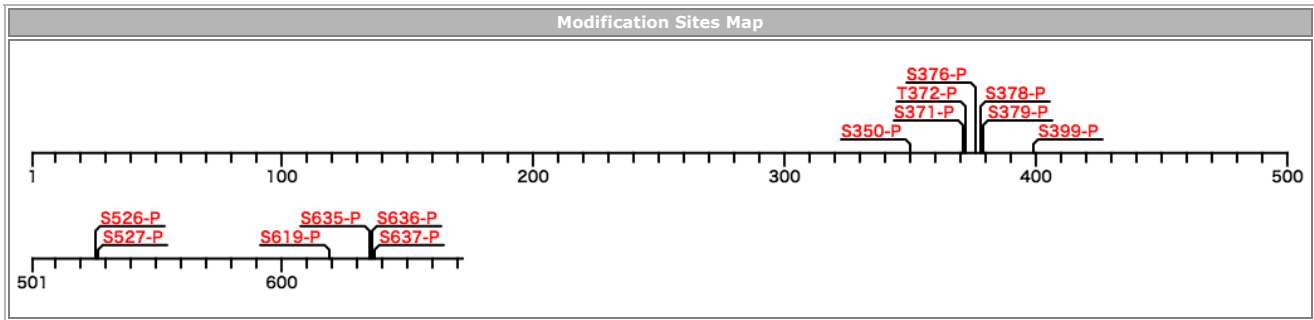


ID	Accession	GeneName	Chr.No.	Description
F169A_HUMAN	Q9Y6X4	FAM169A	5q13.3 74073399..74162776	Soluble lamin-associated protein of 75 kDa



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
350	S	P	Lab	140320_tita_C18_.mgf[F017426]
350	S	P	Lab	140320_tita_SDB_.mgf[F017430]
350	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]

Protein Sequence	
MAFPVDMLEN CSHEELNSA EDYMSDLRCG DPENPECFSL LNITIPISLS NVGFVPLYGG DQTQKILALF APEDSLTAVA LYLADQWWAI DDIVKTSVPS REGLKQVSTL GER VVLYVLN RIIYRKQEME RNEIPFLCHS STDYAKILWK KGEAIGFYSV KPTGSICASF LTQSYQLPVL DTMFLRKKYR GKDFGLHMLE DFVDSFTEDA LGLRYPLSSL MYTACKQ YFE KYPGDHELLW EVEGVGHWYQ RIPVTRALQR EALKILALSQ NEPKRPMSE YGPASVPEYE ARTEDNQSSE MQLTIDSLKD AFASTSEGHK KTSVSTHTRS GNLRPKIG K RFQDSEFSS QGEDEKTSQT SLTASINKLE <b>ST</b> ARPS <b>ESSE</b> EFLEEEPEQR GIEFEDES <b>S</b> D RDARPALETQ PQQEKQDGEK ESELEPMNGE IMDDSLKTSL ITEEEDSTSE VL DEELKLP FNSSEDSTNL VPLVVESSKP PEVDAPDKTP RIPDSEMLMD EGTSDEKGHM EEKLSLLPRK KAHLG <b>SS</b> DNV ATMSNEERSD GGFPNSVIAE FSEEPVSENL SPNTT SSLED QGEEGVSEPEQ ETSTALPQSS LIEVELEDVP FSQNAGQKNQ SEEQSEAS <b>SE</b> QLDQFTQSAE KAVD <b>SSS</b> EIEI EVEVPVDRR NLRKAKGKH GPAKKKAKLT	

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