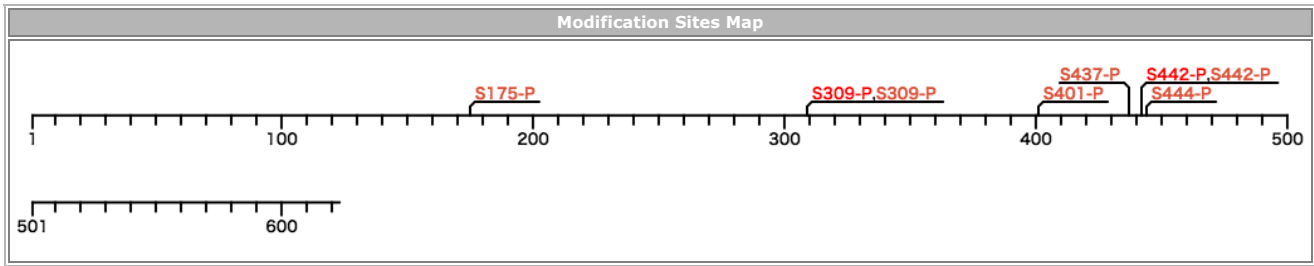


ID	Accession	GeneName	Chr.No.		Description
ELL_HUMAN	P55199	ELL	19p13.11	18553473..18632937	RNA polymerase II elongation factor ELL



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
309	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
309	S	P	Lab	140320_tita_C18_.mgf[F017426]
309	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
309	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
309	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
309	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
309	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MAALKEDRSY GLSCGRVSDG SKVSVFHVKL TDSALRAFES YRARQDSVSL RPSIRFQGSQ GHISIPQPDG PAEARTFSFY LSNIGRDNPQ GSFDCIQQYV SSHGEVHLDC LG SIQDKITV CATDDSYQKA RQSMQAEEEE TRRSIAIVIK AGGRYLGKKV QFRKPAGAT DAVP ^S SRKRAT PINLASAIRK SGASAVSGGS GVSQRPFDRDR VLHLLALRPY RKAEL LLRLQ KDGLTQADKD ALDGLLQVA NMSAKDGTCT LQDCMYKDVQ KDWPGYSEG D QLLKRVLVR KLCQPQSTGS LLGDPAAS ^S P PGERGRSASP PQKRLQPPDF IDPLA NKKPR ISHFTQRAQP AVNGKLGVPN GREALLPTPG PPASTDTLSS STHLPPRLEP PRAHDPLADV ^S NDLGHSGRD CEHGEEAAPA PTVRLGLPLL TDCAQPS ^R RPH GSP ^S RSKP KK KSKKHDKER AAEDKPRAQL PDCAPATHAT PGAPADTPGL NGTCSVSSVP TSTSETPDYL LKYAAISSSE QRQSYKDNFN AEYSEYRDLH ARIERITRRF TQLDAQLRQL SQGSEYETT RGQILQEYRK IKKTNTNYSQ EKHRCEYLHS KLAHIKRLIA EYDQRQLQAW P	

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