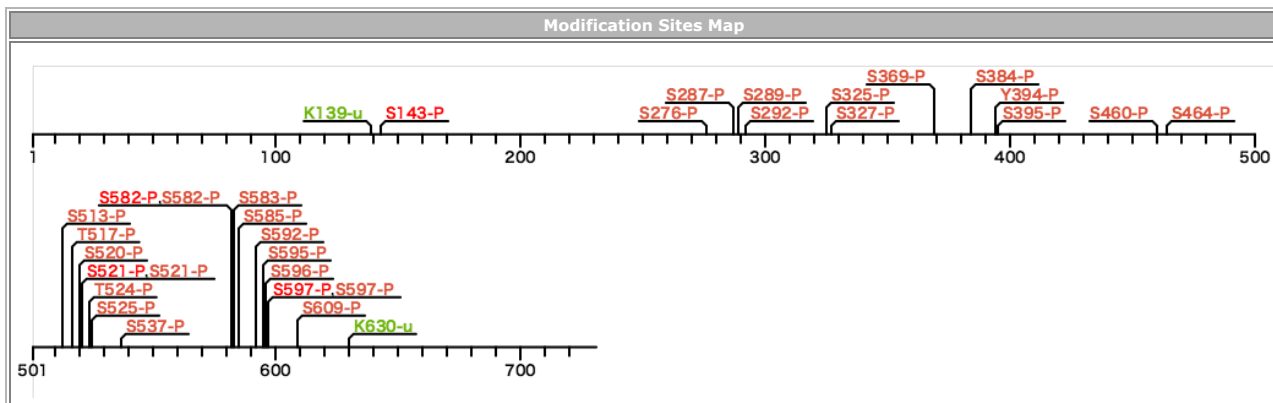


ID	Accession	GeneName	Chr.No.	Description	
BAP1_HUMAN	Q92560	BAP1	3p21.1	52435029..52444366	Ubiquitin carboxyl-terminal hydrolase BAP1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
582	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
582	S	P	Lab	110218_pRMG2_2.mgf[F017476]
582	S	P	Paper	J Proteomics 2014, 96, 253-262

**Protein Sequence**

MNKGWLELES DPGLFTLLVE DFGVKGQVQE EIYDLQSKCQ GPVYGIFILF KWIEERRSRK KVSTLVDDTS VIDDIVNNM FFAHQLIPNS CATHALLSVL LNCSSVDLGP TLS  
 RMKDFTK GFSPEKGYA IGNAPELAKA HNSHARPEPR HLPEKQNGLS AVRTMEAFHF VSYVPITGRL FELDGLKVYP IDHGPWGEDE EWTDKARRVI MERIGLATAG EPYHDI  
 RFNL MAVVPDRRIK YEARLHVLKV NRQTVLEALQ QLIRVTQPEL IQTHKSQESQ LPEESKASAN KSPVLVLEANR APAASEGNHT DGAEAAAGSC AQAPSHSPPN KPCLVVKPP  
 G SSLNGVHPNP TPVQRLPAF LDNHNYAKSP MQEEEDLAAG VGRSRVPVPR PQQYSDDDD YEDDEDDVQ NTNSALRYKG KGTGKPGALS GSADGQLSVL QPNTINVLA  
 E KLKESQKDL S IPLSIKTSSG AGSPAVAVPT HSQPSPTPSN ESTDTASEIG SAFNSPLRSP IRSANPTRPS SPVTSHISKV LFGEDDSL LR VDCIRYNRAV RDLGPIVSTG LLHL  
 AEDGVL SPLALTEGGK GSSPSIRPIQ GSQSSSPVE KEVVEATDSR EKTGMVRPGE PLSGEEKYSPK ELLALLKCV EAEIANYEACL KEEVEKRKKF KIDDQRRTHN YDEFICTF  
 IS MLAQEGMLAN LVEQNISVRR RQGVSIGRLH KQRKPDRRR SRPYAKKR

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