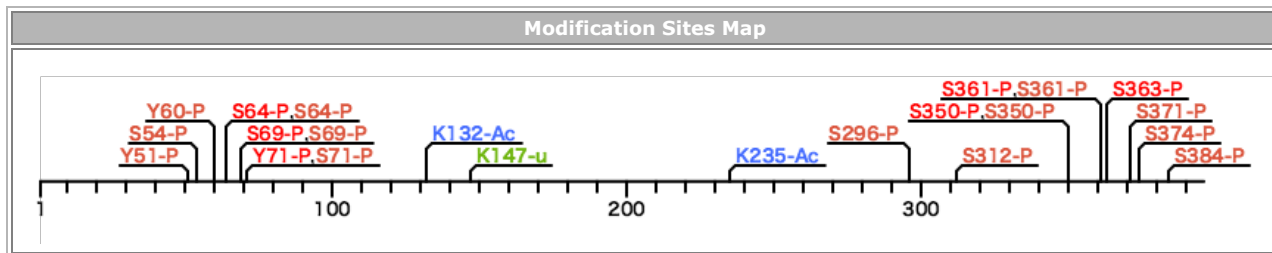


ID	Accession	GeneName	Chr.No.		Description
ALKB5_HUMAN	Q6P6C2	ALKBH5	17p11.2	18086392..18113268	RNA demethylase ALKBH5



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
64	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
64	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
64	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
64	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
64	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
64	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
64	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
64	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
64	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
64	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
64	S	P	Paper	Proc Natl Acad Sci USA 2008, 105(38), 10762-10767
64	S	P	Paper	Sci Signal 2009, 2(84), ra46
64	S	P	Paper	J Proteomics 2014, 96, 253-262

Protein Sequence	
MAAASGYTDL REKLKSMSTR DNYKAGSREA AAAAAA VAA AAAAAAAEP YPVSGAKRKY QEDSDPERSD YEEQQLQKEE EARKVKSGIR Q MRLFSQDEC AKIEARIDEV VSRAEKGLYN EHTVDRAPLR NKYFFGEGYT YGAQLQKRGPGQERLYPPGD VDEIPEWVHQ LVIQKLVEHR VIPE GFVNSA VINDYQGGC IVSHVDPIHI FERPIVSVSF FSDSALCFGK KFQFKPIRVS EPLVLSLPVRR GSVTVLSGYA ADEITHCIRP QDIKERRAVI ILRKTRLDAP RLETKSLSSS VLPPSYASDR LSGNNRDPAL KPKRSHRKAD PDAHRPRIL EMDKEENRRS VLLPThRRRG SFSSSENYWRK SYE SSEDCE AAGSPARKVK MRR	

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