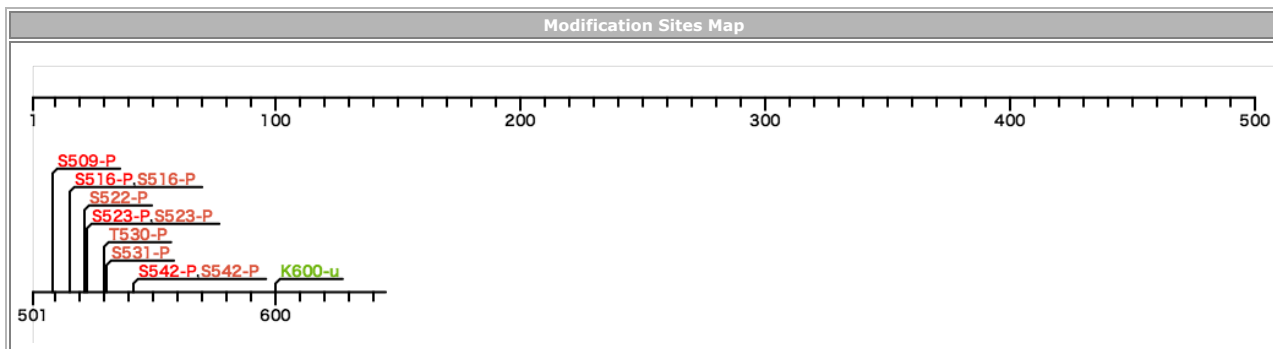


ID	Accession	GeneName	Chr.No.	Description	
AMFR_HUMAN	Q9UKV5	AMFR	16q12.2	56395364..56459450	E3 ubiquitin-protein ligase AMFR



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
516	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
516	S	P	Lab	130626_Gly_S_MGFPeaklist.mgf[F015371]
516	S	P	Paper	Proc Natl Acad Sci USA 2008, 105(38), 10762-10767
516	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence

MPLFLERFP WPSLRTYTGL SGLALLGTII SAYRALSQPE AGPGEPDQLT ASLQPEPPAP ARPSAGGPRA RDVAQYLLSD SLFVWVLVNT ACCVLMVLAK LIQCIVFGPL RVSER QHLKD KFWNFIFYKF IFIFGVLNVQ TVEEVVMWCL WFAGLVFLHL MVQLCKDRFE YLSFSPITPM SSHGRVLSLL VAMLLSCCGL AAVCSITGYT HGMHTLAFMA AESLLVTV RT AHVILRYVIH LWDLNHEGTW EGKGTYYVYT DFMELTLLS LDLMHHIHM LFGNIWLSMA SLVIFMQLRY LFHEVQRRIR RHKNYLRVVG NMEARFVAVT PEELAVNDD CAICWDSMQA ARKLP CGHLF HNSCLRSWLE QDTSCPTCRM SLNIADNNRV REEHQGENLD ENLVPVAAA GRPRLNQHNH FFHFDGSRIA SWLPSFSVEV MHTTNILGIT Q ASNSQLNAM AHQIQEMFPQ VPYHLVLQDL QLTRSVEITT DNILEGRIQV PFPTQRSDS I RPALNS PVER PSSDQEEGET S AQTERVPLD L SPRLEETLD FGEVEPESE VEDFE ARGSR FSKSADERQR MLVQRKDELL QQARKRFLNK S SEDDAASES FLPSEGASSD PVTLRRLMLA AAAERRLQKQ QT

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