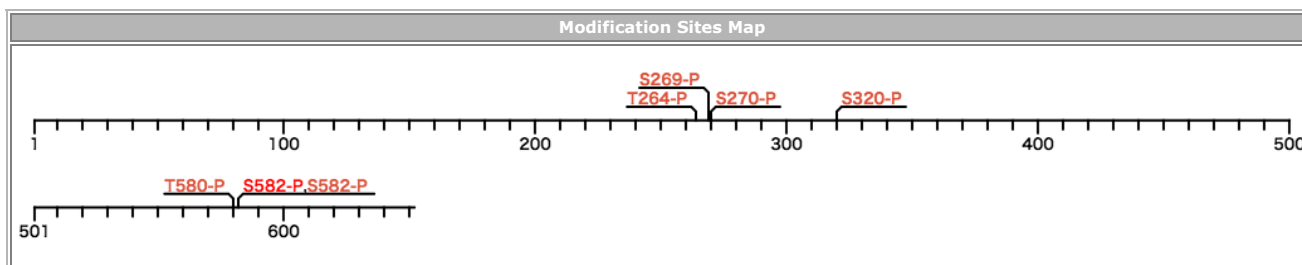


ID	Accession	GeneName	Chr.No.	Description
FAF1_HUMAN	Q9UNN5	FAF1	1p32.3	50905150..51425935 FAS-associated factor 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
582	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
582	S	P	Lab	110218_pRMG1_3.mgf[F017474]
582	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
582	S	P	Paper	Sci Signal 2009, 2(84), ra46
582	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MASNMDREMI LADFQACTGI ENIDEAITLL EQNNWDLVAA INGVIPQENG ILQSEYGGET IPGPAFNPAS HPASAPTSSS SSAFRPVMPMS RQIVERQPRM LDFRVEYRDR NVD VVLEDTC TVGEIKQILE NELQIPVSKM LLKGWKTGDV EDSTVLKSLH LPKNNSLYVL TPDLPPSSS SHAGALQESL NQNFMLIITH REVQREYNLN FSGSSTIQEV KRNVDL TSI PVRHQLWEGW PTSATDDSMC LAESGLSYPC HRLTVGRRSS PAQTREQSEE QITDVHVMVD SDGDDFEDAT EFGVDDGEVF GMASSALRKSPMPMPENAENE GDALLO FTAE FSSRYGDCHP VFFIGSLEAA FQEAFYVKAR DRKLLAIYLH HDESVLTNVF CSQMLCAESI VSYLSQNFIT WAWDLTKDSN RARFLTMCNR HFGSVVAQTI RTQKTDQFP L FLIIMGKRSS NEVLNVIQGN TTVDELMMRL MAAMEIFTAQ QQEDIKDEDE REARENVKRE QDEAYRLSLE ADRAKREAHE REMAEQFRLE QIRKEQEEER EAIRLSLEQA LP PEPKEENA EPVSKLRIRTPSGEFLERRF LASNKLQIVF DfvASKGFPW DEYKLLSTFP RRDVTQLDPN KSLLEVKLFP QETLFLLEAKE	

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