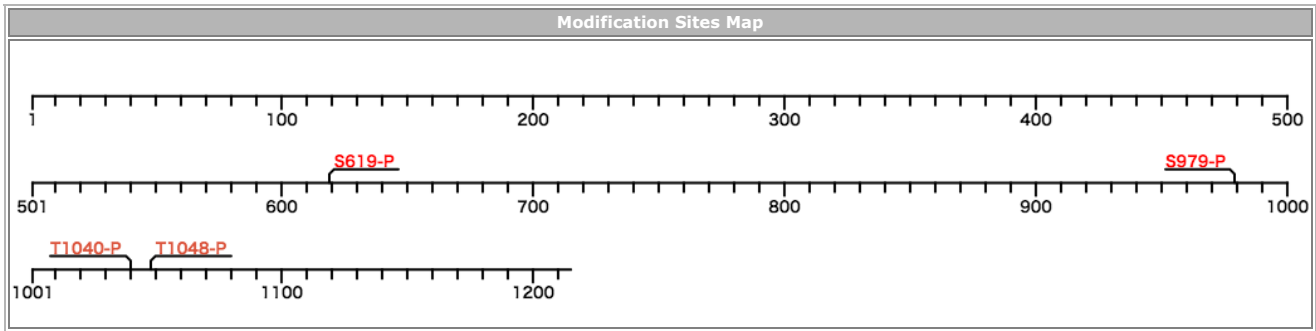


ID	Accession	GeneName	Chr.No.		Description
FLIP1_HUMAN	Q7Z7B0	FILIP1	6q14.1	76001575..76203454	Filamin-A-interacting protein 1



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
979	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
979	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
979	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
979	S	P	Lab	140320_Agarose_.mgf[F017423]
979	S	P	Lab	140320_Agarose_.mgf[F017423]
979	S	P	Lab	140320_OVISE_ME_.mgf[F017429]
979	S	P	Lab	140320_OVISE_ME_.mgf[F017429]
979	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
979	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
979	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
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979	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
979	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
979	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
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979	S	P	Lab	100520-GIST-W3.mgf[F017524]

Protein Sequence
MRSRNQGGES ASDGHISCPK PSIIGNAGEK SLSEDAK KKK KSNRKEDDVM ASGTVKRHLK TSGECERKTK KSELESKEDL IQLLSIMEGE LQAREDVIHM LKTEKTKPEV LE AHYGSAEP EKVLRLVLRD AILAQEKSIG EDVYEKPISE LDRLEEKQKE TYRRMLEQLL LAEKCHRRTV YELENEKHKH TDYMNKSDDF TNLLEQERER LKKLLEQEKA YQARKE KENA KRLNKLRLDEL VKLKSFALML VDERQMHIEQ LGLQSQKVDQ LTQKLREEEE KLKAITSKSK EDQRKLLKLE VDFEHKASRF SQEHEEMNAK LANQESHNRQ LRLKLVGL TQ RIEELEETNK NLQKAEELQ ELRDKIAKGE CGNSSLMAEV ENLRKRVLEM EGKDEEITKT ESQCRELRKK LQEEHHHSKE LRLEVEKLOK RMSELEKLEE AFSKSKSECT Q LHLNLEKEK NLTKDLLNEL EVVKSrvKEL ECSESRLEKA ELSLKDDLTK LKSFTVMLVD ERKNMMEKIK QEERKVDGLN KNFKVEQGVV MDVTEKLEIE SKLLKLLKSE MEEK VYNLTR ERDELIGLKL SEEEKSSELS CSVDLLKKRL DGIEEVEREI TRGRSRKGS E LTCPEDNKIK ELTLEIERLK KRLQQLEVVE GDLMKTEDEY DQLEQKFRTE QDKANFLS QQ LEEIKHQIAK NKAIEKGEVV SQAELRHRP RLEEAKSRDL KAEVQALKEK IHELMNKEDQ LSQQLVDYSV LQQRFMEEEN KNKNMGQEVV NLTKLELSK RYSRALRPSV NGRRMVDVPV TSTGVQTDVAV SGEAAEEETP AVFIRKSFQE ENHIMSNLRQ VGLKPKVERS SVLDYPPAA NELTMRKSWI PWMRKRENGP SITQEKGPRT NSSPGHPGEV V LSPKQGQPL HIRVTPDHN STATLEITSP TSEEFFSSTT VIPTLGNQKP RITIIPSPNV MPQKQKSGDT TLGPERAM SP VTITTFREK TPESGRGAGA DRPTSPIQIM TVSTSA PAE IAVSPESQEM PMGRTILKV I PEKQTVP I PV RKYNSNANII TTEDNKIHH LGSQFKRSPG TSGEGVSPVI TVRPVNVTAE KEVSTGTVLR SPRNHLSSRP GASKVTSTIT IT PVTSSAR GTQSVSGQDG SSQRPTPRI PMSKGMKAGK PVVAAPGAGN LTKFEPRAET QSMKIELKKS AASSTSLGG GKG

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