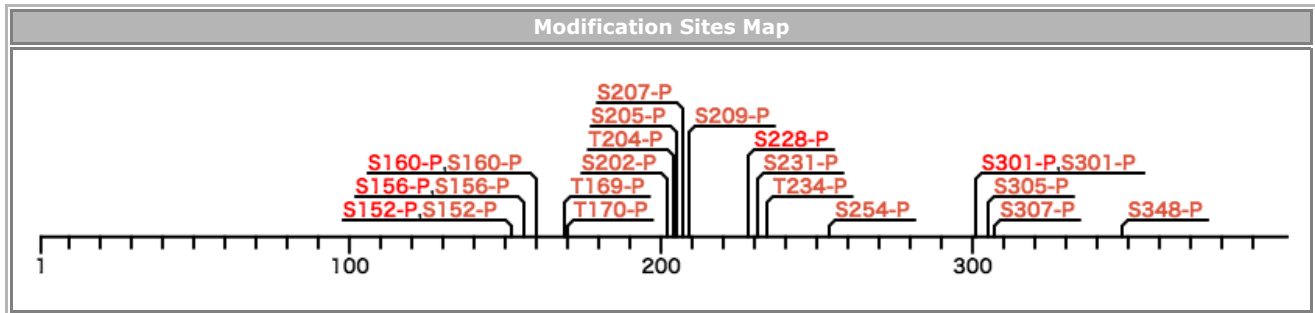


ID	Accession	GeneName	Chr.No.		Description
FR1OP_HUMAN	O95684	FGFR1OP	6q27	167412670..167466201	FGFR1 oncogene partner



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
160	S	P	Lab	130327_HEK_CE_pphos.mgf[F015007]
160	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
160	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
160	S	P	Lab	140320_Agarose_.mgf[F017423]
160	S	P	Lab	140320_tita_C18_.mgf[F017426]
160	S	P	Lab	140320_tita_C18_.mgf[F017426]
160	S	P	Lab	140320_HEK_SCE_.mgf[F017428]
160	S	P	Lab	140320_tita_SDB_.mgf[F017430]
160	S	P	Lab	140320_tita_SDB_.mgf[F017430]
160	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
160	S	P	Paper	Sci Signal 2009, 2(84), ra46
160	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MAATAAAVVA EEDTELRLDLL VQTLNSGVL NRIKAE LRAA VFLALEEQEK VENKTPLVNE SLKKFLNTKD GRLVASLVAE FLQFFNLDFT LA VFQPETST LQGLEGRENL ARDLGIIEAE GTVGGPLLE VIRRCQQKEK GPTTGEGALD L <b>S</b> DVH <b>S</b> PPK <b>S</b> PEGK TSAQ <b>T</b> PSKIPRYK GQ GKK KTS GQKA GDKKANDEAN Q <b>S</b> D <b>T</b> S <b>V</b> S <b>L</b> S E PKSKSSLHLL SHETKIG <b>S</b> FL <b>S</b> NR <b>T</b> LDGKDK AGLCPDEDDM EGD <b>S</b> FFDDPI PKPEKTYGLR KE PRKQAGSL ASLSDAPPLK SGLSSLAGAP <b>S</b> LK D <b>S</b> E <b>S</b> KRG NTVLKD LKLI SDKIGSLGLG TGEDDDYVDD FNSTSHR <b>S</b> EK SEISIGEEIE EDL SVEIDDI NTSDKLDLDT QDLTVS QLS D VADYLEDVA	

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