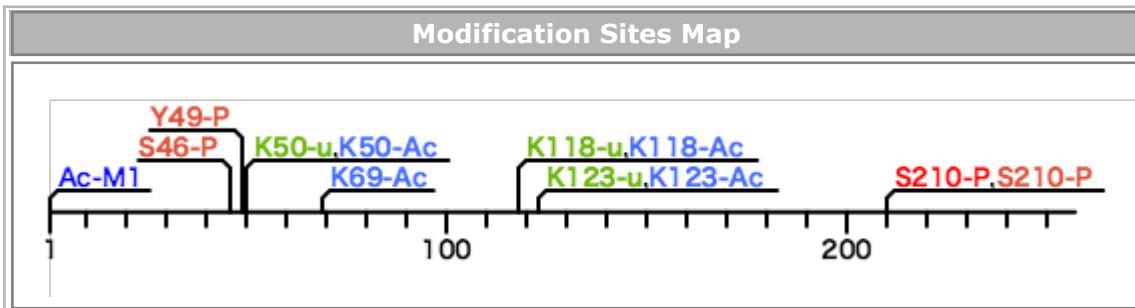


ID	Accession	GeneName	Chr.No.	Description	
1433E_HUMAN	P62258	YWHAE	17p13.3	1247566..1303672	14-3-3 protein epsilon



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
210	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
210	S	P	Paper	Sci Signal 2010, 3(104), ra3

Protein Sequence

MDDREDLVYQ AKLAEQAERY DEMVESMKKV AGMDVELTVE ERNLLSVAYK NVIGARRASW RIISSIE
 QKE ENKGGEDKLIK MIREYRQMVE TELKLICCDI LDVLDKHLIP AANTGESKVF YYKMKGDYHR YLAEF
 ATGND RKEAAENSLV AYKAASDIAM TELPPTHPIR LGLALNFSVF YYEILNSPDR ACRLAKAAFD DAIA
 ELDTLS EESYKDSTLI MQLLRDNLTL WTSDMQGDGE EQNKEALQDV EDEN

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