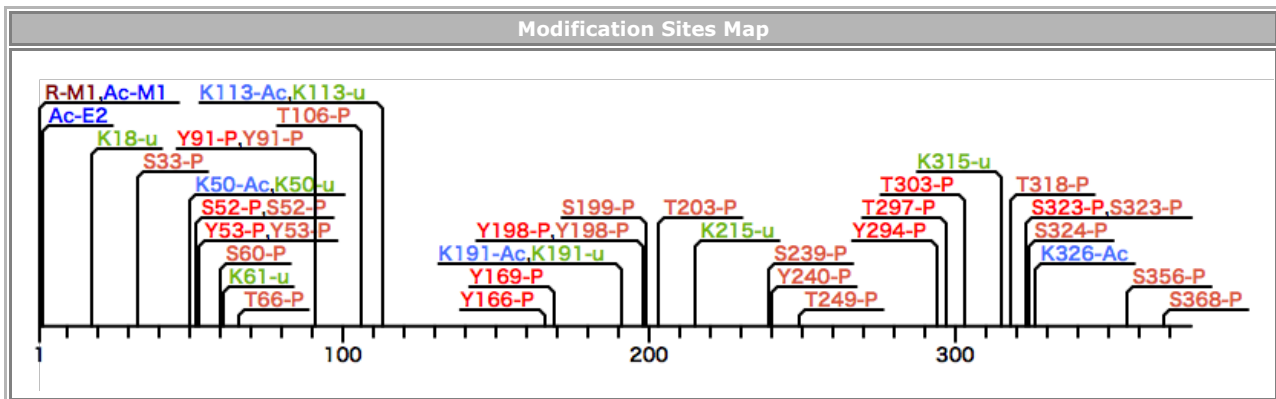


ID	Accession	GeneName	Chr.No.		Description
ACTG_HUMAN	P63261	ACTG1	17q25.3	79476997..79490873	Actin, cytoplasmic 2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
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Protein Sequence

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MEEEIAALVI DNGSGMCKAG FAGDDAPRAV FPSIVGRPRH QGVMVGMGQK DSYVGDEAQS KRGILT LKYP IEHGIVTNWD DMEKIWH
HTF YNELRVAPEE HPVLLTEAPL NPKANREKMT QIMFETFTP AMYVAIQAVL SLYASGRITG IVMDSGDGVV HTVPIYEGYA LPHAILRL
DL AGRDLTDYLM KILTERGYSF TTTAEREIVR DIKEKLCYVA LDPEQEMATA ASSSSLEKSY ELPDGGVIT I GNERFRCPEA LFPSPFLGM
E SCGIHETTFN SIMKCDVDIR KDLYANTVLS GGTIMYPGIA DRMQKEIT AL APSTMKIKII APPERKYSVW IGGISILASLS TFQQM WISK
Q EYDESGPSIV HRKC
    
```

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