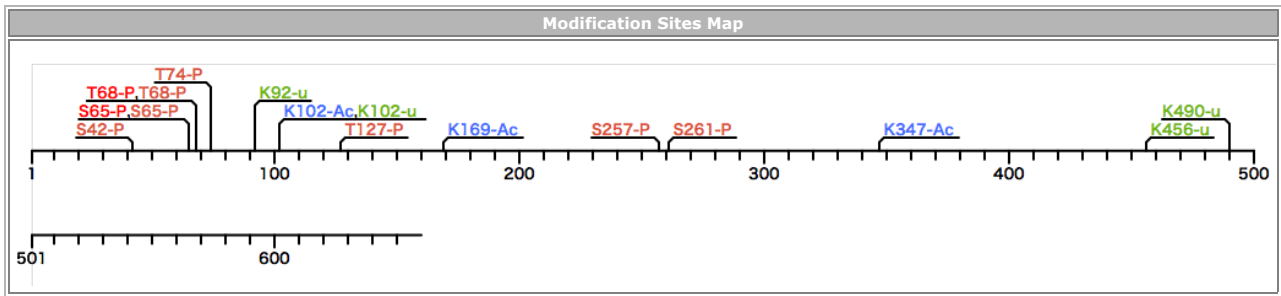


ID	Accession	GeneName	Chr.No.		Description
ADAS_HUMAN	O00116	AGPS	2q31.2	178257372..178408564	Alkyldihydroxyacetonephosphate synthase, peroxisomal



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
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Protein Sequence	
MAEAAAAAGG TGLGAGASYG SAADRDRDPD PDRAGRRLRV LSGHLLGRPR EALSTNECKA RRAASAAATAA PTATPAAQES GTIPKKRQEV MKWNGWGYND SKFIFNKKGQ IELTGKRYPL SGMGLPTFKE WIQNTLGVNV EHKTTSKASL NPSDTPPSVV NEDFLHDLKE TNISYSQEAD DRVFRAHGHC LHEIFLLREG MFERIPDIVL WPTCHDDVVK IVNL ACKYNL CIPIGGGTS VSYGLMCPAD ETRTIISLDT SQMNRILWVD ENNLTAHVEA GITGQELERQ LKESGYCTGH EPDSLEFSTV GGWVSTRASG MKKNIYGNIE DLVVHIK MVT PRGIIEKSCQ GPRMSTGPGDI HHFIMGSEGT LGVITEATIK IRPVPEYQKY GSVAFPNFEQ GVAACLEIAK QRCAPASIRL MDNKQFQFGH ALKPQVSSIF TSFLDGLKKF Y ITKFKGFDP NQLSVATLLF EGDREKVLQH EKQVYDIAAK FGGLAAGEDN GQRGYLLTYV IAYIRDLALE YVVLGESFET SAPWDRVVDL CRNVKERITR ECKEKGVQFA PFSTC RVTQT YDAGACIYFY FAFNYRGISD PLTVFEQTEA AAREEILANG GSLSHHHGVG KLRKQWLKES ISDVGFGMLK SVKEYVDPNN IFGNRNL	

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