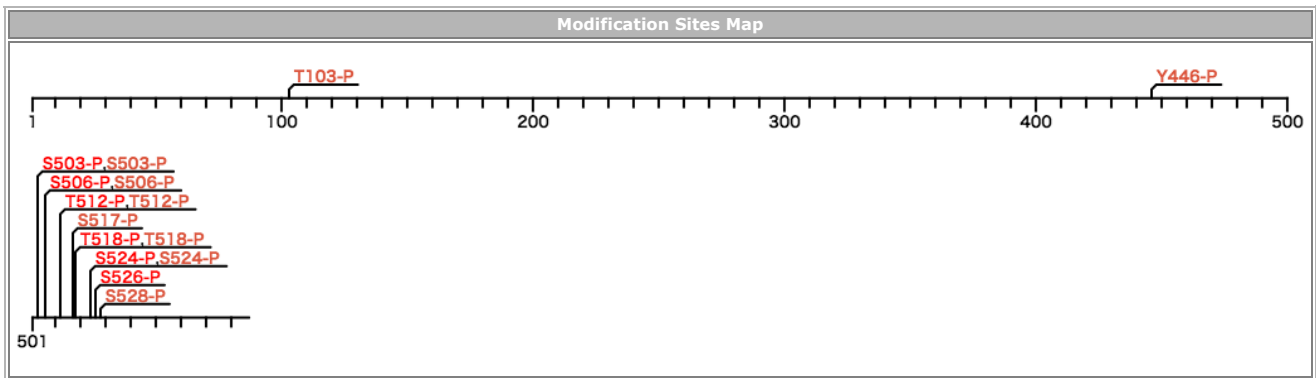


ID	Accession	GeneName	Chr.No.		Description
EIF2A_HUMAN	Q9BY44	EIF2A	3q25.1	150264465..150302029	Eukaryotic translation initiation factor 2A



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
503	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
503	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
503	S	P	Lab	110711_titania_LNcaP_AI_6.mgf[F017448]
503	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
503	S	P	Lab	140320_OVISE_SCE.mgf[F017431]
503	S	P	Lab	100520-GIST-IM1.mgf[F017509]
503	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
503	S	P	Paper	Sci Signal 2009, 2(84), ra46
503	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MAPSTPLLTV RGSEGLYMVN GPPHFTSTV FPRESGKNCK VCIFSKDGTL FAWNGEKNV IISVTNKGLL HSFDLLKAVC LEFSPKNTVL ATWQPYYTTSK DG	YAGIPNLQ LYD
VKTGTCL KSFQKMQN WCPWSWSEDET LCARNVNEV HFFENNNFNT IANKLHLQKI NDFVLSPPGQ PYKVAVYVPG SKGAPSFVRL YQYPNFAAPH AALANKSFFK ADKVT	
MLWNK KATAVLVIAS TDVDKTGASY YGEQTLHYIA TNGESAVVQL PKNGPIYDVV WNSSSTEFCA VYGFMPAKAT IFNLKCDPVF DFGTGPRNAA YYSPHGHLV LAGFGNL	
RGQ MEVWDVKNYK LISKPVASDS TYFAWCPDGE HILTATCAPR LRVNNGYKIW HYTGSIHKY DVPSNAELWQ VSWQPFLDGI FPAKTITYQA VPSEVPNEEP KVATAY	RPP
A LRNKPITNSK LHEEPPQNM KPQSGNDKPL SKTALKNQRK HEAKKAAKQE AR	SDKSPDLA PTPAQSTPR NTV
KNQLEKIQK ETALLQELED LELGI	SISGD PEIDKKIKNL KKKLKAIEQL KEQAATGKQL E

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