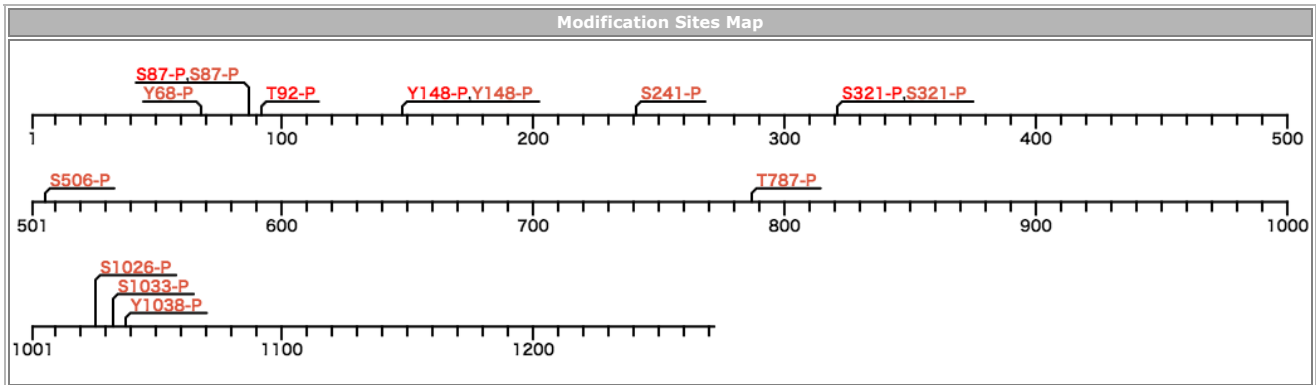


ID	Accession	GeneName	Chr.No.	Description
DHX9_HUMAN	Q08211	DHX9	1q25.3 182808504..182856886	ATP-dependent RNA helicase A



Click a modification site to display the information in detail.

Site no	Amino acid	Type	Division	Detail
87	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
87	S	P	Paper	Cancer Res 2009, 69(6), 2663-2668

Protein Sequence	
MGDVKNFLYA WCGKRKMTPS YEIRAVGNKN RQKFMCEVQV EGYNYTGMGN STNKKDAQSN AARDFVNYLV RINEIKSEEV PAFGVA S PPP L T DTPTTAN AEGDLPTTMG GPLPPLHALK AENNSEVGAS GYGVPGPTWD RGANLKD Y YS RKEEQEVQAT LESEEVDLNA GLHGNWTLEN AKARLNQYFQ KEKIQGEYKY TQVGPDHNS FIAEMTIYIK QL GRRIFARE HGSNKKLAAQ S CALSLVRQL YHLGVVEAYS GLTKKKEGET VEPYKVNLSQ DLEHQLQNI QELNLEILPP PEDPSVVAL NIGKLAQFEP S QRQNQVGVV PWSPP QSNWN PWTSSNIDEG PLAFATPEQI SMDLKNELMY QLEQDHLQA ILQERELLPV KKFESILEA ISQNSVVIIR GATGCGKTTQ VPQFILDFFI QNDRAAECNI VVTQPRRIS A VSAERVAFE RGEPEGKSCG YSVRFESILP RPHASIMFCT VGVLLRLEA GIRG S HVIV DEIHEDINT DFLLVLRDV VQAYPEVRIV LMSATIDTSM FCEYFFNCPI IEVYG RTYPV QEYFLEDCIQ MTHFVPPPKD KKKKDKDDDDG GEDDDANCNL ICGDEYGPET RLMSQLNEK ETPFELIEAL LKYIETLNVP GAVLVFLPGW NLIYTMQKHL EMNPHFGS HR YQILPLHSQI PREEQRKVFDPVPVGVTKVI LSTNIAETSI TINDVVYVID SCKQVKLFT AHNNMTNYAT VWASKTNLEQ RKGRAGRVRP GFCFHLCSRA RFERLE T HMT P EMFRTLHE IALSIKLLRL GGIGQFLAKA IEPPLDAVI EAEHTLRELD ALDANDELTP LGRILAKLPI EPRFGKMMIM GCIFYVGDAI CTIAAATCFP EPFINEGKRL GYIHRNFAG N RFSDHVALLS VFQAWDDARM GGEEAEIRFC EHKRLNMATL RMTWEAKVQL KEILINSQFP EDCLLTQVFT NTGPDNNLDV VISLLAFGVY PNVCYHKEKR KILTTEGRNA L IHK S VNCP FS S QDMK Y PS PFFVFGKIR TRAISAKGMT LVTPLQLLF ASKKVQSDGQ IVLVDDWIKL QISHEAAACI TGLRAAMEAL VVEVTKQPAI ISQLDPVNER MLNMI RQISR PSAAGINLMI GSTRYGDGPR PPKMARYDNG SGYRRGGSSY SGGYGGGYS SGGYGGGYS GSANSFRAGY GAGVGGGYRG VSRGGFRGNS GGDYRGPSGG YR GSGGFQRG GGRGAYGTGY FGQGRGGGGY	

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