

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	er
Q9V4C8	HCF_DROME	Drosophila melanogaster	Host cell factor	32.646721	NaN	S477;S958;S966;T1126;S1489	27918560;33925313;28604694	MEGSDFVDPAFSSGERISADLNSE HIQAEENHSFANRISMDMDVDPGH QLDSNLTGFRWKRVLNPTGPQPRPR HGHRAINIKELMVVFGGNEGIVDE LHVYNTVTNQWYVPLKGDVPNGC AAYGFVVEGTRMFVFGGMIEYGKYS NELYELQATKWEWRKMYPESPDSG LSPCPRLGHSFTMVGEKIFLFGGLA NESDDPKNNIPKYLNDLYILDTRGV HSHNGKWIVPKTYGDSPPRESHTG ISFATKSNGLNLLIYGGMSGCRLG DLWLLTDSMTWSKPKTSGEAPLPR SLHSSTMIGNKMYVFGGWVPLVIND SKSTTEREWKCTNLTAVLDLETMT WENVTLDTVEENVPRARAGHCAVGI QSRLYVWSGRDGYRKAWNNQVCCCK DLWYLEVSKPLYAVKVALVRASTHA LELSWTATTFAAAYVLQIQKIEQPLN TSSKLLSNNIVQQGTPSAETSGINIS ANRSGSALGLGVEATSTVLKLEKESL QLSGCQPETNVQPSVNDLLQSMSQ PSSPASRADKDPLSSGGGTTFNLSTS VASVHPQISVISSTAAVTGNDTASPS GAINSIQKFRPVVTAVRTSTTTAVSI ATSTDPLSVRVPSTMSANVVLSSSS STLRIVPSVTASHSLRIASSQASGNN CRSSSAINILKTALPNVAVQSQPTSS TTTSGGKQYFIQKPLTLAPNVQLQF VKTSGGMTVQTLPKVNFTASKGTPP HGSIANPHLASGITQIQGSTVPGSQI QKPIVSGNVLKLVSPTHMAGGKLM KNSNILQMGKVTNVMGGKPAFVIT NKQGTPLGNQQHIVTTGGNVRVPT STVMTSAGGSASGTNIVSIVNSTSTT PSPLQALSGQKTLISNOSGVKMLRNI SSVQASSMAFGQKQSGTPIHQKTA LYIGGKAVTVMSTNTSMAASGNKV MVLPGTSSNNSPATTALSARKSFV FNAGGSPRTVTLATKSINAKSIPQSQ PVTEINNHSVATIKDTPMDDIIEQL DGAGDLLKLESEGGQHGSEENENN GENATSSSASALFTGGDTAGPSRAO NPIVMEHPVDIIEDVSGVSTTDVNE TAIVSGDTIESLKMSEKENDDVKSM GEKSILSDDCHQPTTSETEAATILTTI KSAEALVLETAEIRKDHTGCTIGSLK ENQDENKFKQRQESSQNIHQF QNVDSQLEALASAALLQAATSDTT ALALKELIERPESETNTRSSNIAEIQQ NNVQSTLAVVPNTSQENQKWHT VGVFKDLSHTVTSYIDSNCSISDFD GIDVNDLPDFSKFPRTNLEPGTAYRF RLSAINSCGRGEWGEISSFKTCLPGF PGAPSAIKISKDVKEGAHLTWEPPPA QKTEIIEYSVYLAVKPTAKDKALSTP QLAFVRVYVGAANQCTVFNASLSNA HVDCSNKPAIFRIAARNQKGYGPAT QVRWLQDPAAAKQHTPTVTPNLKR GPEKSTIGSSNIANTFCSPHKGRGN GLHD	None	None	None	None	None	None	None	N	