

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7TMY8	HUWE1_MOUSE	Mus musculus	E3 ubiquitin-protein ligase HUWE1	19.294551	S2619;S2632	S648;S649;S740;S1084;S1368;S1370;S1382;S1395;S1907;T2035;S2266;S2362;S2365;S2391;S2527;S2532;S2535;T2554;S2584;S2595;S2619;T2751;S2826;S2833;S2835;S2861;S2887;S2888;T2889;S2918;S3116;S3117;S3122;S3127;S3135;S3557;S3663;S3753;S3758;S3760;S3761;S3810;S3818;S3830;T3833;S3909;S3922;T3927;T3930;T4274	36288343;29125738;36852467;33300544	<p>MKVDRTKLLKKTPTPEAPADCRALIDK LKVCNDEQLLLELQIKTWNIGKCE LYHWVDLLDRFDGILADAGQTVEN MSWMLVCDRPEKEQLKMLLLAVLN FTALLIEYSFSRHLVSSIEHLTLLAS SDMQVVLAVLNLLVFSKRSNYITRL GSDKRTPLLTRLQHLAESWGGKEN FGFLAECCRDLOMLKYPPSATTLHF EFYADPGAEVKIEKRRTSNTLHYIHI EQLDKISESPSEIMESLTKMYSIPKD KQMLLFTHIRLAHGFSNHRKRLQAV QARLHAISILVYSNALQESANSILYN GLIEELVDVLQITDKQLMEIKAASLR TLTSIVHLERTPKLSSIIDCTGTASYH GFLPVLVRNCIQAMIDPSMDPYPHQ FATALFSFLYHLASYDAGGEALVSCG MMEALLKVIKFLGDEQDQITFVTRA VRVVDLITNLDMAAFQSHSGLSIFIY RLEHEVDLCRKECPFVIKPKIQRPT TQEGEEMETDMDGVQCIPQRAALL KSMLNFKKAIQDPAFSDGIRHVMD GSLPTSLKHIIISNAEYYGPSLFLATE VVTVFVFQEPSLLSSLQDNGLTDVM LHALLIKDVPATREVLGSLPNVFSAL CLNARGLQSFVQCQPFERLQVLLS PDYLPAMRRRRSSDPLGDTASNLS AVDELMRHQPTLKTDTATTAIKLEE ICNLGRDPKYICQKPSIQKADGTATA PPRSNHAAEEASSEDEEEEEVQAM QSFNSAQNETEPNQVVGTEERIP IPLMDYILNVMKFVESILSNNTTDD HCQEFVNQKGLLPLVTILGLPNLPID FPTSAACQAVAGVCKSILTSHEPKV LQEGLLQLDLILSSLEPLHRPIESPG GSVLLRELACAGNVADATLSAQATP LLHALTAAHAYIMMFVHTCRVQSE IRSISVNQWGSQGLSVLSKLSQLYC SLVWESTVLLSLCTPNLPSGCEFG QADMQKLVKDEKAGTTQGGKRS GEQDGTAGSMDASAQGLLEGIELDG DTLAPMETDEPSSSDSKGKSKITPA MAARIKQIKPLLSASSRLGRALAEFL GLLVKLCVGSPPVRRRSHHAASSTT APTPAARSTASALTKLLTKGLSWQPP PYTPTPRFRLTFFICSVGFTSPMLFD ERKYPYHMLQKFLCSGGHNAFET FNWALSMGGKVPVSEGLEHSDLPD GTGEFLDAWLMLVEKMNPTTVLE SPHSLPAKLPGGVQSFQFSALRFLV VTQKAAFTCIKNLWNRKPLKVYGG MAESMLAILCHILRGEPVIRERLSKE KEGSRGEEEGQEEGSRREPQVN QQQLQQLMDMGFTREHAMEALLN TSTMEQATEYLLTHPPPIIGGVVRDL SMSEEDQMMRAIAMS LGQDIPMDQ RAESPEEVACRKEEEERKAREKQEE</p>

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