	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q8C1D8	IWS1_MOUSE	Mus musculus	Protein IWS1 homolog	25.685855	NaN	S27;S54;S69;S157;S170;S172;S183;S196;S198;S209;S211;S222;S224;S235;S237;S248;S250;S252;S261;S263;S277;S280;S295;S306;S307;S309;S321;S343;S345;S366;S368;S372;T381;S384;S386;S408;S410;S412;S427;T436;S458;S460;T672	39627609	MDSEYYSGDQSDDGGATPVQDERD SGSDGEDGVTEQHSGSDTGSVDHH SENETSDREDGLAKIHNGTDSENDE PSNANASDSESEELHRPKDSDSDSE EHAESPASDSENEPVNQHGSDSEN EELLNGHASDSEKEEVSKHAASDSE AEDTLQPQVSESDSEDPPRPQASDS ENEEPPKPRISDSESEELPKPRVSDS ESEDPPRPQASDSESEELPKPRVSDS ESEDPPRPQASDSESEELPKPRVSDS ESEDPQKGPASDSEAEDASRHKEKP DSDDSDGENKREDSEVQNESDGHT DRKGLHSSDSEEEEPKRQKIDSDDD EEKEGDEEKVAKRKAAVLSDSEDDA GNASAKKSRVVCDADDSDSDVVSDK SGKRETTVASDSEEEAGKEESSVKK KEDKDLFGSDSESGNEEENLIADIF GESGDEEEEEFTGFNQEDLEEEKNE TQLKEAEDSDSDDNIKRGKHMDFLS DFEMMLQRKKSMCGKRRRNRDGG TFISDADDVVSAMIVKMNEAAEEDR QLNNQKKPALKKLTLLPTVVMHLKK QDLKETFIDSGVMSAIKEWLSPLPD RSLPALKIREELLKILQELPSVSQETL KHSGIGRAVMYLYKHPKESRSNKD MAGKLINEWSRPIFGLTSNYKGMTR EEREQRDLEQMPQRRRLSSTGGQT PRRDLEKVLTGEEKALRPGDPGFCA RARVPMPSNKDYVVRPKWNVEMES SRFQASSKKGISRLDKQMRKFTDIR KKSRSAHAVKISIEGNKMPL	True	False	False	4.706	False	False	False	False	False