

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q920B9	SP16H_MOUSE	Mus musculus	FACT complex subunit SPT16	25.685855	NaN	S188;S455;S508;S650;S658;T903;S979;S982;S986;S1015	39627609	MAVTLDKDAYRRRVKRLYSNWRKG EDEYASIDAIVSVGVDEEIVYAKSTA LQTWLFGEYELTDTIMVFCDDKIIFMA SKKKVEFLKQIANTKGNENANGAPA ITLLVREKNESSKSSFDKMDAIKES KSGKKIGVFSKDKFPGEFMKSWSDC LNKEGFQKVDISAVVYATIAVKEDGE LNLMMKKAASITSEVFNKFFKERVME IVDADEKVRHRSKLAESVEKAIEEKY LAGADPSTVEMCYPPIIQSGGNYNLK FSVSDKNMHMFGAITCAMGIRFKS YCSNVRTLMVDPTQEVQENYNFLL QLQEELLKELRHGVKICDVYNSVMD VVKKQKPELLNKITKNLGFQGMGIEF REGSLVINSKNQYKLLKGMVFSINL GFSDLTNKEGKKPEEKTYALFIGDTV LVDEDGPATILTSVKKVKNVGIFLK NEDDEEEEEKDEAEDLLGRGSRAA LLTERTRNEMTAEKRRRAHQKELAA QLNEEAKRRLTEQKGEQQIQKARKS NVSYKNPSPMPKPHIREMKIYIDKK YETVIMPVFGIATPFHIATIKNISMV EGDYTYLRINFYCPGSALGRNEGNIF PNPEATFVKEITYRASNMKAPGEQT VPALNLQNAFRIKEVQKRYKTREAE EKEKEGIVKQDSLVINLNRSNPKLK DLYIRPNIAQKRMQGSLEAHVNGFR FTSVRGDKVDILYNNIKHALFQPCD GEMIIVLHFHLKNAVMFGKKRHTDV QFYTEVGEITDLGKHQMHDRDD LYAEQMEREMRHLKLTAFKNFIEKV EALTKEELEFEVPPRDLGFNGAPYR STCLLOPTSSALVNATEWPPFVVTLD EVELIHFERVQFHLKNFDMVIVYKD YSKKVTMINAIPVASLDPIKEWLNCS DLKYTEGVQSLNWKIMKTIIVDDPE GFFEQGGWSFLEPEGEGSDAEDGD SESEIEDETFNPSEDDYEEEDSDE DYSSEAEESDYSKESLGSEEEESGKD WDELEEEARKADRESRYEEEEEQSR SMSRKRKASVHSSCRGSRNRSRHS SAPPKKRKR	True	False	2.42	5.0	1.14	0.983	False	0.776	0.713