

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6A065	CE170_MOUSE	Mus musculus	Centrosomal protein of 170 kDa	21.509147	NaN	S141;S355;S358;T363;S443;S463;S494;T498;S568;S577;S628;S631;T639;S662;S718;T752;S829;S870;S872;T906;T912;S922;S925;S950;S1008;T1012;T1047;S1048;S1102;S1104;S1122;S1123;S1135;S1150;S1155;S1188;S1195;S1200;S1229;S1231;S1241;S1260;S1270;S1362;S1521;S1522	34678516;33300544;22645316	MSLTSWFLVSSGGTRHRLPREMIFVGRDDCELMLQSRVSKQHAVINYDASMDEHLVKDLGSLNGTFVNDVRIP EQTYITLKLEDKLRFGYDTNLFVVRGEMRVPEEALKHEKFTIQLQLSQKS SESELPKSASAKGTDSKVEAAAEEVQP RATEALKSEEKPMDEVSAAMPRGTPLY GQPSWWGDAEEDEQRAFKANGKPE GKSQEAGASGCSTEAKHVEGQSAASA SEEALFPFCREPSYFEIPTKEFQQPS QIAESTIHEIPTKDTSSHTAGAGHA SFTIEFDDSTPGKVTIRDHVTKFTSD QRHKSCKASPQTQDLPGIQTGMMA PENKVADWLAQNNPPQMWERTE EDSKSIKSDVPVYLKRLKGNKHDDG TQSDSENAGAHRRCSKRATLEEHLR RHHSEQKKKAQSTEKHQEQAATSST HHRGGHGVPHGKLLKQKSEEPSVS LPFLQTALLRSSGSLGHRPSQEMDV MLKNQATSASSEKDNDDDQSDKGT YTIELENPNSEEEVEARKMIDKVFV DDNQDYNRPINEKHKGLIKDWALN SAAVMEERKPLSTPGFHNSEEAIS SSGSKRWVSQWASLAANHTRHDPE ERLMELSATVENETDTGDAGVSLRS TSCTTSLASQGERKRRTLPLPNEE KLESSRAKVVPQRSEIGEQQDTLQ EKEAQVYQSEKHDADRGLSKMSRA VNGESPKTGGDGKALLHSGSSSKE KSETEKETS LVKQTLAKMQQEQKE QAQWTPTKFPSKNALGHIDKCREES SKQESQLEKVS GHSTSKGDRVIQN ESKRRKAEIIPKQASKGDKKESK SLVRQGSFTIDKPSSNIPIELIPHINK QNSSVPTALALTSASRLRERSDSLDT DSSMDTTLILKDTEAVMAFLEAKLR EDNKNKTDEGPDTPSYNRDNSISPES DVDTASTISLVTGETERKSTQKRKSF TSLYKDRCSTSSPSKDVTKSGSREKI EKKAKSRSADIGARADGRKFVQSSG RIRQPSIDLTDQDQTSVPHSAISDI MSSDQETYSCKSHGRTPLTSADEHN IHSKLEGGKATKSKTSPVAGSTSKS TTLPRPRPRTSLRRARLGEASDSE LADADKASVASEVSTTSSTSKPPTGR RTISRIDLLAQPRRRLGSLARS DS EATISRSSASARTAEAVIRSGARLVPS DKLSRPRTRANSISRLSDSKVKMSST HGSPSVNSRWRRFPTDYASTSEDEF GSNRNSPKHTRLRTSPALKTTRMQS TGSAMPASSSFKHRIKEQEDYIRDW TAHREEIARISQDLALIAREINDVAGE IDSVTSSGTAPSTTVSTAATTPGSAID

TR	EE	VG	DL	HG	EM	HK	LV	DR	VF	DE	SL	NFR	KIP	PL	VH	SK	TP	EG	NN	GR	SV	DS	R	PQ	AE	HP	DL	HT	IR	RR	TW	SR	DE	VM	G	DN	LL	SS	VF	QF	SR	KI	RQ	SID	KTAG	KI	RIL	FK	DK	DR	NW	DD	IE	NK	LR	AE	SE	VPI	VKT	SS	ME	ISS	IL	QEL	KR	VE	KQL	QVIN	AM	ID	PD	GT	LE	AL	NN	MG	FP	NAIL	PSP	PK	QK	SS	PV	NN	HSS	PS	QTP	AL	CP	PET	RAL	HP	AA	AG	VAAA	ASTE	FENA	ESEA	DF	SI	HF	NR	FN	PD	GE	EED	VTV	HE
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