

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region	
0	Q60625	ICAM5_MOUSE	Mus musculus	Intercellular adhesion molecule 5	24.474581	NaN	T182	33300544	MPGSPGLRRALLGLWAALGLGILGI SAVALEPFWADLQPRVALVERGGSL WLNCSSTNCPRPERRGGLETSLRRNG TQRGLRWLARQLVDIREPETQPVCF FRCARRTLQARGLIRTFQRPDRVELV PLPSWQPVGENFTLSCRVPGAGPRA SLTLTLRGGQELIRRSFVGEPPRAR GAMLTARVLARREDHRVNFSCLAEL DLRPHGLGLFANSSAPRQLRTFAMP PHSPSLIAPRVLEVDSERPVTCITLDG LFPAPEAGVYLSLGDQRLNPNVTLD GDSLVAATATASAEQEGTKQLMCV VTLGGESRETOENLTVYSFPTPLTL SEPEAPEGKMTISCWAGARALVTL EGIPAAVPGQPAELQNLVTKNDDKR GFFCDAALDVGGETLRKNQSSSELRV LYAPRLDDLDPCRSWTWPEGPEQTL HCEARGNPEPSVHCARPEGGAVLAL GLLGPVTRALAGTYRCTAVNGGQQA VKDVTTLTVEYAPALDSVGCPEHITWL EGTEASLSCVAHGVPSPVSCVRSR KEEVMEGPLRVAREHAGTYRCEAIN ARGSAAKNVAVTVEYGPSFEELGCP SNWTWVEGSGKLFSCSEVDGKPEPR VECVGSEGASEGIVLPLVSSNSGPR NSMTPGNLSPGIYLCNATNRHGSTV KTVVVS AESPPQMD ESSCP SHQ TWL EGAEATALACSARGRPSRVHCSRE GAARLERLQVSREDAGTYRCVATNA HGTDSRTVTVGVEYRPVVAELAASPP SVRPGGNFTLTCRAEAWPPAQISWR APPGALNGLSSNNSTLSVAGAMGS HGGEYCAATNAHGRHARRITVRVA GPWLWVAVGGAAGGAALLAAGAGL AFYVQSTACKKGEYNVQEAESSGEA VCLNGAGGTPGAEGGAETPGTAESP ADGEVFAIQLTSS	False	True	1.471	1.673	1.457	1.172	1.14	5.0	2.945