	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	Q9UHX3	AGRE2_HUMAN	Homo sapiens	Adhesion G protein-coupled receptor E2	26.21016	T361;S364	NaN	29351928	MGGRVFLVFLAFCVWLTLPGAETQD SRGCARWCPQDSSCVNATACRCNP GFSSFSEIITTPMETCDDINECATLS KVSCGKFSDCWNTEGSYDCVCSPGY EPVSGAKTFKNESENTCQDVDECQ QNPRLCKSYGTCVNTLGSYTCQCLP GFKLKPEDPKLCTDVNECTSGQNPC HSSTHCLNNVGSYQCRCRPGWQPIP GSPNGPNNTVCEDVDECSSGQHQC DSSTVCFNTVGSYSCRCRPGWKPRH GIPNNQKDTVCEDMTFSTWTPPPGV HSQTLSRFFDKVQDLGRDYKPGLAN NTIQSILQALDELLEAPGDLETLPRL QQHCVASHLLDGLEDVLRGLSKNLS NGLLNFSYPAGTELSLEVQKQVDRS VTLRQNQAVMQLDWNQAQKSGDP GPSVVGLVSIPGMGKLLAEAPLVLEP EKQMLLHETHQGLLQDGSPILLSDV ISAFLSNNDTQNLSSPVTFTFSHRSV IPRQKVLCVFWEHGQNGCGHWATT GCSTIGTRDTSTICRCTHLSSFAVLM AHYDVQEEDPVLTVITYMGLSVSLLC LLLAALTFLLCKAIQNTSTSLHLQLS LCLFLAHLLFLVAIDQTGHKVLCSIIA GTLHYLYLATLTWMLLEALYLFLTAR NLTVVNYSSINRFMKKLMFPVGYGV PAVTVAISAASRPHLYGTPSRCWLQP EKGFIWGFLGPVCAIFSVNLVLFLVT LWILKNRLSSLNSEVSTLRNTRMLA FKATAQLFILGCTWCLGILQVGPAAR VMAYLFTIINSLQGVFIFLVYCLLSQQ VREQYGKWSKGIRKLKTESEMHTLS SSAKADTSKPSTVN	False	True	1.581	1.961	1.382	1.343	1.119	5.0	2.215