	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	Q8C419	GP158_MOUSE	Mus musculus	Probable G-protein coupled receptor 158	24.474581	NaN	S694;S705;S708;S865;S944;S1059;S1074	33300544	MGAMAYSLLFCLLLAHLGLGEVGAS LDPPGRPDSPRERTPRGKQHGQQLP RASAPDPSIPWSRSTDGTILAQKLAE EVPVDVASYLYTGDFHQLKRANCSG RYELAGLPGKSPSLASSHPSLHGAL DTLTHATNFLNMMLQSNKSREQTV QDDLQWYQALVRSLLEGEPSISRAAI TFSTESLSTPAPQVFLQATREESRILL QDLSSSAHHLANATLETEWFHGLR RKWRPHLHRRGSNQGPRGLGHSW RRRDGLGGDRSHVKWSPPYLECEN GSYKPGWLVTLSAAFYGLQPNLVPE FRGVMKVDISLQKVDIDQCSSDGWF SGTHKCHLNNSECMPIKGLGFVLGA YQCICKAGFYHPRVFSVNNFQRRGP DHHFSGSTKDVSEETHVCLPCREGC PFCADDRPCFVQEDKYLRLAIISFQA LCMLLDFVSMLVVYHFRKAKSIRAS GLILLETILFGSLLLYFPVVILYFEPST FRCILLRWARLLGFATVYGTVTLKLH RVLKVFLSRTAQRIPYMTGGRVMRM LAVIVLVVFWFLVGWTSSMCQNLER DILLVGQGQTSDHLTFNMCLIDRWD YMTAVAEFLFLLWGIYLCYAVRTVPS AFHEPRYMAVAVHNELIITAIFHTIR FVLASRLQPDWMLMLYFAHAHLTV TVTIGLLLIPKFSHSSNNPRDDIATEA YEDELDMGRSGSYLNSSINSAWSEH SLDPEDIRDELKKLYAQLEIYKRKKM ITNNPHLQKKRCSKKGLGRSIMRRIT EIPETVSRQCSKEDKEGTDHSAAKG TGLVRKNPTESSGNTGRPKEESLKN RVFSLKKSHSTYDHVRDQTVESSSL PMESQEEEATENSTLESLSSKKLTQ KLKEDSEAESTESVPLVCKSASAHN LSSEKKPGHPRTSMLQKSLSVIASAK EKTLGLAGKTQTLVMEDRAKSQKPK DRETIRKYSNSDNVETIPNSGHMEE PRKPQKSGIMKQQRVSLPTANPDVS SGITQIKDNFDIGEVCPWEVYDLTPG PMPSEPKAQKHVSIAASEVEQNPAS FLKEKSYHKSKATEGLYQANHKSID KTEVCPWEIHSQSLLEDENRLISKTP VLPGRAREENGSQLYTTNMCAGQY EELPHKVVAPKVENENLNQMGDQE KQTSSSVDIIPGSCNSSNNSHQPLTS RAEVCPWEFEPLEQPNAERSVTLPA SSALSANKIPGPQK	None	None	None	None	None	None	None	None	None