

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma mem
Q8WXG9	AGRV1_HUMAN	Homo sapiens	Adhesion G-protein coupled receptor V1	29.387686	S2054;T2057;S2327;T2331	NaN	35008409;38253038;29351928	MSVFLGPGMPSASLLVNLLSALLLIFVFGETEIRFTQOTFEVFNSTTTVIRLIIRIGEPANVAIVSLYGEDAGDFFDYAAAFIPAGETNRTVYIACVDDDLPEPDETFIFHLTLQKPSANVKGWPRTVTVTILSNDNAFGIISFNMLPSIAVSEPKGRNESMPLTLIREKGTGMVMVTFEVEGGPNPDEDLSPVKNITFPGRATVIYNLTVLDDEVPEDEIFLIQLKSVEGGAEINTSRNSIEIHKKNDSPVRFLQSIYLVPEEDHILIPVVRGKDNNGNLIGSDEYEVSYAVTTGNSTAHAQQNLDLFDLQPNNTTVVFPFIHESHLKQFQVDDTIPEIAESFHIMLLKDTLQGDVAVLISPSVVQVTIKPNDKPYGVSFNLSVLFERTVIIDEDRISRYEITVVRNGGTHGNVSANWVLRNSTDPSPTADIRPSSGVLHFAQQQMLATIPLTVVDDDLPEEAEAYLLQILPHTIRGGAEVSEPAELLFYIQDSDDVYGLITFFPMENQKIESSPGERYLSLFFTRLGGTKGDVRLLYSVLYPAGAVDPLQAKEGILNISRRNDLIFPEQKTQVTTKLPINRDAFLQNGAHFLVQLTEVELLNIIPLIPPISPRFGIEICNISLLVTPAIANGEIGFLSNLPIILHEPEDFAAEVYIPLHRDGTGQATVYVSLKPSGFNSKAVTPDDIGPFGNSVFLSGQSDTTINITIKGDDIPEMNETVTLSDRVNVENQVLKSGYTSRDLIILENDPFGGVFEFSPASRGYVVIKESGESVELHIIIRSGSLVKQFLHYRVEPRDSNEFYGNTGVLEFKPGEREIVITLLARLDGPELDEHYVWVLSHGERESKLGSAITVNITLKNDDPHGIIIEFVSDGLIVMINESKGDAYSAVYDVVRNRGNFGDVSWSVVSPDFTQDVFPVQGTVVFGDQEFKNIITYSLPDEIPEEMEEFTVILLNGTGGAKVGNRTTATLRIRRNDDPIYFAEPRVVRVQEGETANFTVLRNGSVDVTCMVQYATKDGKATARERDFIPVEKGETLIFEVGSROQSSISIFVNEDGIPETDEPFYILLNSTGDTVYQYGVATVIEANDDPNGIFSLPIDKAVEEGKTNFVILRHRGYFGSVSVWQLFQNDLALQPGQEFYETSGTVNFMDEGEEAKPILHAFPDKIPEFNEFYFLKLVNISGSGPGGQLAETNLQVTVMVFPNDDPFGVFILDPECLEREVAEDVLSDEDDMSYITNFTILRQQGVFGDVQLGWEILSSEFPAGLPPMIDFLLVGIFPTTVHLQQHMRRHHSRNTDIALYFTGLEGAFGTVPKYPYHSRNTTIANFTFSAWVMPNANTNGFIKAKDDGNGSIYGVKIQTNESHVTLSLHYKTLGNSATYIAKTTVMKYLEESVWLHLLIILEDGIIIEFYLDGNAMPRGISLKGAEITDGPILRIGAGINGNDRFTGLMQDVRSYERKLTLEIYELHAMPAKSDLHPISGYLEFRQGETNKSFIISARDNDDEEGEELFILKLVSVYGGARISEENTTARLTIQKSDNANGLFGFTGACIPEIAEEGSTISCVVERTRGALDYVHVFTYISQIETDGINYLVDDFANASGTITFLPWORSEVLNIVLDDDIPELNEYFRVTLVSAIPGDGKLGSTPTSGASIDPEKETTDITIKASDHPYGLLQFSTGLPPQPKDAMTLPASSVPHITVEEDGEIRLLVIRAQGLLGRVTAEFRTVSLTAFSPEDYQNVACTLEFQPGERYKYIFINITDNSIPELEKSFVLELLNLEGGVAELFRVDGSGSGDGMMEFFLPTIHKRASLGVASQILVTAASDHAHGVFEFSPESLFSVSGTEPEDGYSTVTLNVIRHHGTLSPVTLHWNIDSDPDGLAFTSGNITFEIGQTSANITVEILPDEDPELDKAFSVSVLSVSSGSLGAHINATLTVLASDDPYGIFIFSEKNRPVKVEATQNITLSIIRLKGMLGKVLVSYATLDDMEKPPYFPPNLRATQGRDYIPASGFAFLGANQSEATIAISILDDDEPERSESV	False	True	1.302	1.696	1.469	1.152	0.697	4.726

FIELLNSTLVAKVQSRSPNSPRLGP
 KVETIAQLIHANDDAFGTLQLSAPIV
 RVAENHVGPIHNVTRTGGAFADVSVK
 FKAVPITAJAGEDYSIASSDVVLEGE
 TSKAVPIYVINDIYPEEESFLVQLM
 NETTGGARLGALTEAVIIIASDDPY
 GLFGFQITKLIVEEPEFNsvkVNLPII
 RNSGTLGNVTVQVWATINGQLATGD
 LRVVSGNVTFAPGETIQTLLEVLAD
 DVPEIEEVIQVQLTDAASGGGTIGLDRI
 ANIIPANDDPYGTVAFAQMVYRVQE
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 QGVDPDLWRTWMNVSAVGEPLYTC
 ATCLCKEQACSAFSFFSASEGPQCF
 WMTSWISPAVNNSDFWYRKNMTR
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 QEGDEFANLTVSILPDDFPEMDES
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 LQFHVIRTFPGRGNVTVNWKIIQON
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