

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q9P2B2	FPRP_HUMAN	Homo sapiens	Prostaglandin F2 receptor negative regulator	23.693478	S394;S395	T271	28510447;38253038	MGRLASRPLLLALLSLALCRGRVVR VPTATLVRVGTTELVIPCNVSDYDGP SEQNFDFWSFSSSLGSSFVELASTWEV GFPAQLYQERLQRGEILLRRTANDA VELHIKNVQPSDQGHYKCSPTSDA TVQGNVEDTVQVKVLADSLHVGPSA RPPPSLSLREGEPEFELRCTAASASPL HTHLALLWEVHRGPARRSVLALTHE GRFHPGLGYEQRYHSGDVRDLDTVGS DAYRLSVSRALSADQGSYRCIVSEWI AEQGNWQEIQEKAVEVATVVIQPSV LRAAVPKNVVAEGKELDLTCNITTD RADDVRPEVTWSFSRMPDSTLPGSR VLARLDRDSLVSHPHVALSHVDAR SYHLLVRDVSKEENSGYYCHVSLWA PGHNRSWHKVAEAVSSPAGVGVTV LEPDYQVYLNASKVPGFADDPTELA CRVVDTKSGEANVRFTVSWYYRMN RRSDNVVTSSELLAVMDGDWTLKYG ERSKQRAQDGFIFISKEHTDTFNFR IORTTEEDRGNYCVVSAWTKORN SWVKSQDVFSKPVNIFWALEDVSLV VKARQPKPFFAAGNTFEMTKVSSK NIKSPRYSVLMAEKPVGDLSSPNET KYIISLDQDSVVKLENWTDASRV VLEKQVEDEFYRMYQTQVSDAGL YRCMVTAWSPVVRGSLWREAATSLS NPIEIDFQTSGPVFNASVHSDTPSVIR GDLIKLFCIITVEGAALDPDDMAFDV SWFAVHSFGLDKAPVLLSSDRKGI VTTSTRRDWKS DLSLERVSVLEFLLO VHGSEDQDFGNYYCSVTPWVKSPTG SWQKEAEIHSKPVFITVKMDVLNAF KYPLLIGVGLSTVIGLLSCLIGYCSSH WCCKKEVQETRRERRRLMSMEMD	False	False	1.39	1.373	1.078	3.474	3.442	3.386	2.743