

| 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 |
|--------------|------------|--------------|--|--------------|---------------|-----------------------|----------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| Q9WV91 | FPRP_MOUSE | Mus musculus | Prostaglandin F2 receptor negative regulator | 24.474581 | NaN | T271 | 33300544 | MGRFPAPRPLLALLSLAVCRGRVVR VPAGTLVRRVVGTELVIPCNVSDYDGP SEQNFDWSFSSSGSSFVELASTWEV GFPAQLYRERLQRGDILLRRTANDA VELHIKNVQPSDQGHYKCSTPSTDA TVQGNIEDTVQVKVLADALVVGPS RPPPGLSLREGEFPFELRCIASTTSPL HTHLALRWELHRGPVHRSILALSHE GRFHPPGPGYEQRYHSGDVRDLTVGS DAYRLSVARALSADQGSYRCVVSEW ITEQGSWQEIQEKAWEVATVVIQPTA LQLAVPRTVSVTEGKDLDLSCNITTD RVDDVRPEVTWYFKKTPDTSLASH MLARLDRDSL VHSSPHVALSHVDTR SYHLLVRDVSKENSGYYLCLVALWA PGHNRSWHKVAEAMSAPSGVSVTW LEPEYQVYLNASKVPGFSDDPTELQ CRVIDTKRLEAGVRLTVSWYRMTR RNDDVVASELLAVMDGDWTLRYGE RSKQRAQDGEFIFSKEHTDTFNFR IRTTEEDRGNYCVVSAWTRQRNN SWVKSADVFSKPVNIFWASEDSVLV VKARQPKPFAAGNTFEMTCKVSSK NIKSPRYSVLITAEKPVGDLSNET KYIISLDQDSVVKLENWTDASRVDG VVLEKVQEDEFYRMYQTQVSDAGL YRCMVTAWSPIGGSLWREAATLSLN PIEIDFQTSPTFNASVHSDTPSVTR GDLIKLFCIVTEGAVLDPDDMAFDV SWFAVHSFGLDKAPVLLSSDRKGV VTTGQRDVKSTVSLERVSVLEFLQ VHGSEDQDFGNYYCSVTPWVRSPT GSWQREAEIHSRPIFIVKMDVINA FKYPLLIQVGLSTVIGLLSCLIGYCSS HWCCCKEVRETRRRRLMSMEM D | False | False | 1.335 | 1.307 | 1.02 | 3.47 | 3.342 | 3.301 | 1.841 |