| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatu |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | Q32MZ4 | LRRF1_HUMAN | Homo sapiens | Leucine-rich repeat flightless-interacting protein 1 | 15.050987 | T477 | S16;S115;S116;S120;S124;S555;S564;S581;S618;S638;T676;S714;S733;S766;S768 | 30379171;38665916;35132862;23301498;26853435;28510447;29351928 | MTSPAAAQSREIDCLSPEAQKLAEA<br>RLAAKRAARAEAREIRMKELERQQK<br>EEDSERYSRRSRRNTSASDEDERMS<br>VGSRGSLRVEERPEKDFTEKGSRNM<br>PGLSAATLASLGGTSSRRGSGDTSISI<br>DTEASIREIKELNELKDQIQDVEGKY<br>MQGLKEMKDSLAEVEEKYKKAMVS<br>NAQLDNEKTNFMYQVDTLKDMLLE<br>LEEQLAESRRQYEEKNKEFEREKHA<br>HSILQFQFAEVKEALKQREEMLEKH<br>GIILNSEIATNGETSDTLNNVGYQGP<br>TKMTKEELNALKSTGDGTLGRASEV<br>EVKNEIVANVGKREILHNTEKEQHT<br>EDTVKDCVDIEVFPAGENTEDQKSS<br>EDTAPFLGTLAGATYEEQVQSQILES<br>SSLPENTVQVESNEVMGAPDDRTRT<br>PLEPSNCWSDLDGGNHTENVGEAA<br>VTQVEEQAGTVASCPLGHSDDTVYH<br>DDKCMVEVPQELETSTGHSLEKEFT<br>NQEAAEPKEVPAHSTEVGRDHNEE<br>EGEETGLRDEKPIKTEVPGSPAGTEG<br>NCQEATGPSTVDTQNEPLDMKEPD<br>EEKSDQQGEALDSSQKKTKNKKKK<br>NKKKKSPVPVETLKDVKKELTYQNT<br>DLSEIKEEEQVKSTDRKSAVEAQNE<br>VTENPKQKIAAESSENVDCPENPKIK<br>LDGKLDQEGDDVQTAAEEVLADGDT<br>LDFEDDTVQSSGPRAGGEELDEGVA<br>KDNAKIDGATQSSPAEPKSEDADRC<br>TLPEHESPSQDISDACEAESTERCE<br>MSEHPSQTVRKALDSNSLENDDLSA<br>PGREPGHFNPESREDTRGGNEKGK<br>SKEDCTMS | None | None | None | None | None | None | None |