

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|------------------------------------|--------------|---------------|--|----------------------------|--|
| Q9Y2X7 | GIT1_HUMAN | Homo sapiens | ARF GTPase-activating protein GIT1 | 11.532908 | S561;S575 | T224;S359;S362;T364;S370;S375;T383;S385;S388;T392;S410;S413;S417;T480;S498;S536;T537;T545;T554;S561;S571;S592;S596;T601;S630 | 35132862;34019948;28657654 | MSRKGPRAEVCADCSAPDPGWASIS RGVLVCDECCSVHRSLGRHISIVKH LRHSAWPPTLLQMVHTLASNGANSI WEHSLLDPAQVQSGRRKANPQDKV HPIKSEFIRAKYQMLAFVHKLPCR DDGVTAKDLKQLHSSVRTGNLETC LRLSLGAQANFFHPEKGTTPHVA AKAGQTLQAE LLVY GADPGSPDVN GRTPIDYARQAGHHELAERLVECQY ELTDRLAFYLCGRKPDHKN GHYIIPQ MADSLDLSELAKAAKKLQALSNRL FEELAMDVYDEVDRRENDVWLAT QNHSTLVTERS AVPFLPVNPEYSAT RNQGRQKLARFNAREFATLIIDILSE AKRRQQGKSLSSPTDNLELSLRSQS DLDDQHDYDSVASDEDTDQEPLRST GATRSNRARSMDSDDLSDGAVTLQE YLELKKALATSEAKVQQLMKVNSSL SDELRRLLQREIHKLQAENLQLRQPP GPVPTPLPSEAEHTP MAPGGSTH RRDRQAFSMYEPGSALKPFGPPGD ELTTRLQPFHSTELEDDAIYSVHVPA GLYRIRKGV SASAVPFTPSSPLLS QEGSRHTSKLSRHGSGADSDYENT QSGDPLLGLEGRFLELGKEEDFHP ELES LDGDLDPGLPSTEDVILKTEQV TKNIQELLRAAQEFKHDSFVPCSEKI HLAVTEMASLFPKRPALEPVRSSLRL LNASAYRLQSECRKTPPEPGAPVD FQLLTQQVIQ CAYDIAKAAKQLVTITT REKKQ |