

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|------------------|--------------|---------------|-----------------------|----------------------------|--|
| P23229 | ITA6_HUMAN | Homo sapiens | Integrin alpha-6 | 5.945205 | S452 | S1064;S1059;S1103 | 33214551;33465208;34019948 | MAAAGQLCLLYLSAGLLSRLGAAFN LDTREDNVIRKYGDPSLFGFSLAM HWQLQPEDKRLLLVGAPRAEALPLQ RANRTGGLYSCDITARGPCTRIEFDN DADPTSESKEDQWMGVTVQSQGGP GKVVTCAHRYEKROHVNTKQESRDI FGRCYVLSQNLRIEDDMDGGDWSF CDGRLRGHEKFGSCQQGVAATFTK DFHYIVFGAPGTYNWKGIVRVEQKN NTFFDMNIFEDGPYEVGGETEHDES LVPVPANSYLGLLFLTSVSYTDPDQF VYKTRPPREQPDTFPDVMMSYLGF SLDSGKGIKDEITFVSGAPRANHS GAVLLKRDMSAHLLEPHIFDGE LASSFGYDVAVVDLNKDGWQDIVIG APQYFDRDGEVGGAVVYMNQQGR WNNVKPIRLNGTKDSMFGIAVKNIG DINQDGYPDIAVGAPYDDLKGVFIYH GSANGINTKPTQVLKGISPYFGYSIA GNMDLDRNSYPDVAVGSLSDSVTIF RSRPVINIQTITVTPNRIDLKQKTAC GAPSGICLQVKSCFEYTANPAGYNPS ISIVGTLEAEKERRKSGLSSRVQFRN QGSEPKYTQELTLKRQKQKVCMEET LWLQDNIRDKLRPIPTASVEIQEPSS RRRVNSLPEVLPILNSDEPKTAHIDV HFLKEGCGDDNVCNSNLKLEYKFC TREGNQDKFSYLPKQGVPELVKLD QKDIALEITVTNSPSNPRNPTKDGDD AHEAKLIATFPDTLTYSAYRELRAFP EKQLSCVANQNGSQADCELGNPFK RNSNVTFYLVLTTEVTFDTPDLIN LKLETTSNQDNLAPITAKAKVVI LSVSGVAKPSQVYFGGTVVGEQAMK SEDEVGSLIEYFRVINLGKPLTNLG TATLNIQWPKEISNGKWLKLVKVE SKGLEKVTCEPQKEINSLNLTESHN SRKKREITEKQIDNRKFSLFAERKY QTLNCSVNVNVCNIRCPRLGLDSKA SLILRSRLWNSTFLEEYSKLNLDIL MRAFIDVTAANIRLPNAGTQVRV TVFPSKTVAQYSGVPWWILVAILAGI LMLALLVFILWKCFFKRSRYDDSV PRYHAVRIRKEEREIKDEKYIDNLEK KQWITKWNENESYS |