

| 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 | ext reg |
|--------------|-------------|--------------|-------------------------------------------------|--------------|---------------|-----------------------|----------------------------------------------------------------------------------|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|---------|
| 000232 | PSD12_HUMAN | Homo sapiens | 26S proteasome non-ATPase regulatory subunit 12 | 18.248065 | NaN | NaN | 23301498;32870666;38665916;34725712;40914422;35008409;34019948;35138101;28510447 | MADGGSERADGRIVKMEVDYSATV DQRLPECAKLAKEGRLQEVETLLSL EKQTRTASDMVSTSRILVAVVKMCY EAKEWDLLENIMLLSKRRSOLKQA VAKMVQCCTYVEITDLPKLRID TLRMVTEGKIYVEIERARLTKLATIK EQNGDVKEAASILQELQVETYGSME KKERVEFILEQMRLCLAVKDYIRTQII SKKINTKFFQEENTEKLKLYYNLMI QLDQHEGSYLSICKHYRAIYDTPCIQ AESEKWQALKSVVLYVILAPFDNE QSDLVHRISGDKKLEIIPKYDLLKL FTTMELMRWSTLVEDYGMELRKGS LESPATDVFSGTEEGEKRWKDLKNR VVEHNIRIMAKYYTRITMKRMAQLL DLSVDESEAFLSNLVVKTIFAKVDR LAGHNFQRPKDPNNLLNDWSQKLN SLMSLVNKTTHLIAKEEMIHNLQ | True | False | 4.734 | 4.643 | 1.765 | 1.55 | 0.957 | 1.213 | 4.25 |