

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
P29590	PML_HUMAN	Homo sapiens	Protein PML	13.878804	NaN	S8;S36;S38;S48;S117;S403;S493;S504;S505;S512;S518;S527;S530;S565;T867;S565;S518;S527;S530	34019948;33214551;23301498;38665916	MEPAPARSPRPQDPARPQPTMPP PETPSEGRQPSPPSPTERAPASEEE FQFLRCQQCAEAKCPKLLPCLHLL CSGCLASGMQCPICQAPWPLGADT PALDNVFFESLQRRRLSVYRQVDAQA VCTRCKESADFWCFECEQLLCAKCF EAHQWFLKHEARPLAELRNQSVRE FLDGTTRKTNNIFCSNPNHRTPTLTSI YCRGCSKPLCCSCALLDSSHSELKC DISAEIQRQEELDAMTQALQEQDS AFGAVHAQMHAAVGQLGRARAETE ELIRERVQVVAVHRAQERELLEAV DARYORDYEEMASRLGRLDAVLORI RTGSALVQRMKCYASDQEVLDMHG FLRQALCRLRQEEPOSQAQAVRTDG FDEFKVRQLDLSSCITQGKDAAVSK KASPEAASTPRDPIDVDLPEEAERVK AQVQALGLAEAQPMVAVVQSVPGAHP VPVYAFSFKGSPSYGEDVSNITTAQKR KCSQTQCPKVKIKMESEEGKEARLA RSSPEQPRPSTSKAVSPPHLDGPPSP RSPVIGSEVFLPNSNHVAVSGAGEAE ERVVVISSSESDAENSSSRELDSS SESSDLQLEGPSTLRVLDENLADPQ AEDRPLVFFDLKIDNETQKISQLAAV NRESKFRVVIQPEAFFSIYSKAVSLE VGLQHFLSFLSSMRRRILACYKLWG PGLPNFFRALEDINRLWFEQAEISGF LAALPLIRERVPGASSFKLNLAQTY LARNMSERSAMAAVLAMRDLCRLL EVSPGPQLAQHVYPFSSLQCFASLQ PLVQAAVLPRAEARLLALHNVSFME LLSAHRRDRQGGKYSRYLSLQTT TLPPAQPAFNLQALGTYFEGLLEGPA LARAEGVSTPLAGRGLAERASQQS	True	False	4.396	5.0	1.99	3.695	1.33	1.80