

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasm meml
P22102	PUR2_HUMAN	Homo sapiens	Trifunctional purine biosynthetic protein adenosine-3	20.144112	T39;T706	S10;S440;T682;S796;S802	30397120;23301498;33214551;40307207;32870666;16408927;38665916;34725712;32119511;40914422;35132862;39302247;34019948;35138101;28510447	MAARVLIIGSGGREHTLAWKLAQSH HVKQVLVAPGNAGTACSEKISNTAIS ISDHTALAQFCKEKKIEFVVVGP LAAGIVGNLRSAGVQCFCGPTAEAAQ LESSKRFAKEFMDRHGIPTAQWKA TKPEEACSFILSADFPALVVKASGLA AGKGVIVAKSKEEACKAVQEIMQEK AFGAAGETIVIEELLDGEEVSLCFT DGKTVAPMPPAQDHRKRLLEGGGP NTGGMGAYCPAPQVSNLKKIKDT VLQRTVDGMQQEGTPTGILYAGIM LTKNGPKVLEFNCRFGDPECCQVILP LLKSDLYEVIQSTLDGLLCTSLPVWL ENHTALTVMASKGYPGDYTKGVEI TGFPEAQAQGLEVFHAGTALKNGKV VTHGGRVLAVTAIRENLISALEEAKK GLAAIKFEGAIYKDVGFRAIAFLQQ PRSLTYKESGVDAAGNMLVKKIQPL AKATRSRGCKVDLGGFAGLFDLKAA GFKDPLLASGTDGVTGLKIAQLCN KHDTIGQDLVAMCVNDILAQGAELP FFLDYFSCGKLDLSVTEAVVAGIACA CGKAGCALLGGETAEMPDMYPPGE YDLAGFAVGAMERDQKPLHLERITE GDVVVGIASSGLHNSGFSVLRKIVAK SSLQYSSPAPDGCQDQTLGDLTPT RIYSHSLLPVLRSRSHVKAFAHITGGG LLENIPRVLPEKLGVDLDAQTWIRPR VFSWLQEQEGHLSSEEMARTFNCV GAVLVVSKEQTEQILRDIQHKKEA WVIGSVVARAEGSPRVKVNLI QINGSVLKNGLTNHFSFEKKARV AVLISGTGSLQALIDSTREPNSAQ IDIVISNKAAVAGLDKAERAGIPTRVI NHKLYKNRVEFDSALDLVLEFSIDI VCLAGFMRLSGPFVQKWNKMLNI HPSLLPSFKGSNAHEQALETGVTVT GCTVHFVAEDVDAGQIILQEAVPVKR GDTVATLSERVKLAEHKIFPAALQLV ASGTVQLGENGKICWVKEE	True	False	5.0	3.257	3.459	1.707	0.855	1.975