

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	extracellular region
P35228	NOS2_HUMAN	Homo sapiens	Nitric oxide synthase, inducible	18.379595	NaN	S234;T575;S578;S892	30379171	MACPWKFLFKTKFHQYAMNGEKDI NNNVEKAPCATSSPVTQDDLQYHN LSKQQNESPPQLVETGKKSPESLVK LDATPLSSPRHVRKINWGSMTFQD TLHHKAKGILTCRSKSLGSIPTPKS LTRGPRDKPTPPDELLPQAEFVNQY YGSFKEAKIEEHLARVEAVTKEIETT GTYQLTGDELIFATKQAWRNAPRCI GRIQWSNLQVFDARSCSTAREMFE HICRHVRYSTNNGNIRSAITVFPQRS DGKHDVFRVWNAQLIRYAGYQMPDG SIRGDPANVEFTQLCIDLGWPKPYG RFDVVPVLVQANGRDPPELFEIPDLV LEVAMEHPKYEWFRELELKWYALPA VANMLLEVGGLEFPGCPFNGWYMG TEIGVRDFCDVQRYNILEEVGRRMG LETHKLASLWKDQAVVEINIAVLHSF QKQNVTIMDHHSAAESFMKYMONE YRSRGGCPADWIWLVPPMSGITPV FHQEMLNIVLSPFYQQVEAWKTH VWQDEKRRPKRREIPLKVLVKAFLF ACMLMRKTMASRVRTILFATETGK SEALAWDLGALFSCAFNPKVVCMD KYRLSCLLEERLLVVTSTFCNGDCP GNGEKLLKSLFMLKELNNKFRYAVF GLGSSMYPRFAFAHDIDQKLSHLG ASQLTPMGEDELSGQEDAFRSWA VQTFKAACETFDVRGKQHIQIPKLYT SNVTWDPHHYRLVQDSQPLDLSKAL SSMHAKNVFTMRLKSRQNLQSPST SRATILVELSCEGGQLNYLPGHEHL GVCPCGNQPALVQGILERVVDGPTPH QTVRLEALDESGSYWVSDKRLPPCS LSQALTYFLDITTPPTQLLQKLAQV ATEEPERQRLEALCQPSEYSKWKFT NSPTFLVLEEFPSLRVSAAGFLLSQL PILKPRFYSISSRDHTPTEIHLTVAV VTYHTRDGGPLHHGVCSTWLNLSL KPQDPVPCFVRNASGFHLPEDPSHP CILIGPGTGIAPFRSFQQRLHDSQH KGVRRGGRMTLVFGCRRPDEDHIYQE EMLEMAQKGVLHAVHTAYSRLPGK PKVYVQDILRQQLASEVLRVLHKEPG HLYVCGDVRMARDVAHTLKQLVAA KLKLNEEQVEDYFFQLKSQKRYHED IFGAVFPYEAKKDRVAVQPSLEMSA L	True	False	4.437	4.725	3.186	2.327	1.527	4.121	2.99

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