

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q969R2	OSBP2_HUMAN	Homo sapiens	Oxysterol-binding protein 2	21.366884	NaN	S287;S763	28510447	MGKAAAPSRGGGCGGRSRLSSLFT VVPCLSCHTAAPGMSASTSGSGPEP KQPQPVPPEPERGPLSEQVSEAVSE AVPRSEPVSETTSEPEPGAGQPSELL QGSRPGSESSSGVGAGPFTKAASEP LSRAVGSATFLRPESGSLPALKPLPL LRPGQAKTPLGVPMSGTGTSSAPL ALLPLDSFEGWLLKWTNYLKGQRR WVFLGNLLSYRNRQGEAHTCRG TINLSTAHIDTEDSCGILLTSGARSY HLKASSEVDRQQWITALELAKAKAV RVMNTHSDDSGDDDEATTPADKSE LHHTLKNLSLKLDDLSTCNDLIAKH GAALQRSLTELDGLKIPSESGEKLKV VNERATLFRITSNAMINACRDFLELA EIHSRKWQRALQYEQEQRVHLEETI EQLAKQHNSLERAFHSAPGRPANPS KSFIEGSLTPKGEDSEDEDETEYFD AMEDSTSFITVITEAKEDSRKAEGST GTSSVDWSSADNVLDGASLVPKGSS KVKRRVRIPNKNPNYSLNLWSIMKNC IGRELSRIPMPVNFNEPLSMLQRLT EDLEYHLLDKAVHCTSSVEQMCL VAAFSVSSYSTTVHRIAKPFNPMLGE TFELDRLDDMGLRSLCEQVSHHPPS AAHYVFSKHGWSLWQEITISSKFRG KYISIMPLGAIHLEFQASGNHYVWR KSTSTVHNIIVGKLWIDQSGDIEVN HKTNDRCQLKFLPYSYFSKEAARKV TGVVSDSQGAHYVLSGSWDEQME CSKVMHSSPSSSDGKQKTVYQTL SAKLLWKYPLPENAEENMYFSELA LTLNEHEEGVAPTD SRLRPDQRLME KGRWDEANTEKQRLLEEKQRLSRRR RLEACGPGSSCSSEEEKEADAYTPL WFEKRLDPLTGEMACVYKGGYWEA KEKQDWHMCPNIF	True	False	4.369	1.983	2.275	3.786	1.612	3.544	1.742