

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
Q13873	BMPR2_HUMAN	Homo sapiens	Bone morphogenetic protein receptor type-2	27.086674	T805	T379;S586;S680;S681	28657654	MTSSLQRPWRVPWLPWTILLVSTAA ASQNQERLCAFKDPYQODLGIGESR ISHENGTILCSKSGSTCYGLWEKSKG DINLVKQGCWWSHIGDPQECHYEECV VTTTPPSIQNGTYRFCCSSTDLCNV NFTENFPPDITPLSPPHSFNRETI ILALASVSVLAVLIVALCFGYRMLTGD RKQGLHSMNMMEAAASEPSLDLD NLKLELIGRGRYGAVYKGLDERPV AVKVFSFANRQNFINEKNIYRPLM EHDNIARFIVGDERVTADGRMEYLL VMEYYPNGSLCKYLSLHTSDWVSSC RLAHSVTRGLAYLHTELPGRGDHYKP AISHRDLNSRNVLVKNDGTCVISDF GLSMRLTGNRLVVRPGEEDNAAISEV GTIRYMAPEVLEGAVNLRDCEALK QVDMYALGLIYWEIFMRCTDLFPGE SVPEYQMAFQTEVGNHPTFEDMQV LVSREKQRPKFPEAWKENSLAVRSL KETIEDCWDQDAEARLTAQCAEER MAELMMIWERNKSVPTVNPMTA MQNERNLNRRRVPKIGYPDYSSS SYIEDSIHHTDSIVKNISSEHMSST PLTIGEKNRNSINERQQAQARIPSP ETSVTSLSTNTTTNTTGLTPSTGMT TISEMPYPDETNLHTTNVAQSIGPTP VCLQLTEEDLETNKLDPKEVDKNLK ESSDENLMEHSLKQFSGPDLSSSTS SSLLYPLIKLAVEATGQDFTQTANG QACLIPDVLPTQIYPLPKQNLKRP TSLPLNTKNSTKEPRLKFGSKHKSN LKQVETGVAKMNTINAAEPHVVTVT MNGVAGRNVSVNSHAATTQYANGT VLSGQTTNIVTHRAQEMLQNFQFIGE DTRLNINSSPDEHEPLLRREQQAGH DEGVLDRLVDRRERPLEGGRTNSN NNNSNPCSEQDVLQGVPTAADP GPSKPRRAQRPNSLDLSATNVLDGS SIQIGESTQDGKSGSGEKIKRVKTP YSLKRWRPSTWVISTESLDCEVNNN GSNRAVHKSSTAVYLAEGGTATTM VSKDIGMNCL	None	None	None	None	None	None	None	None	

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