

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrio
Q15910	EZH2_HUMAN	Homo sapiens	Histone-lysine N-methyltransferase EZH2	50.434381	S73;S75;S84;S87;T313;S729	S21;S76;T339;T345;S363;S366;T367;T487	29941599;38561350;30093632;38494860;35470239;24474760;34510715	MGQTGKSEKGPVCWRKRVKSEYM RLRQLKRFRRADEVKSFMSSNRQKI LERTEILNQEWKQRRIQPVHILTSVS SLRGTRCSVTSDDLDFPTQVIPLKTL NAVASVPIMYSWSPLQQNFMVEDE TVLHNIPYMGDEVLDQDGTFFIEELIK NYDGKVVHGDRECCGFINDIEFVELVN ALGQYNDDDDDDDGDDPEEREKQ KDLEDHRDDKESRPPRFPSDKIFE AISSMFPDKGTAEEELKEYKELTEQ QLPGALPPECTPNIDGPNKSVQRE QSLHSFHTLFCRRCFKYDCFLHPFH ATPNTYKRKNTETALDNKPCGPQCY QHLEGAKEFAAALTAERIKTPPKRPG GRRRCRLPNNSSRPSTPTINVLESK DTSDREAGTETGGENNDKEEEEK KDETSSESSEANSRCQTPKMKPNIEP PENVEWSGAEASMFRLIGTYDDNF CAIARLIGTKTCRQVYEFVKESSIIA PAPAEDVDTPPRKKRKHRLWAAHC RKIQLKKDGSNNHVNYQPCDHPRQ PCDSSPCVIAQNFCEKFCQCSSEC QNRFPGCRCKAQCNTKQCPCYLAV RECDPDLCLTCGAADHWDSKNVSC KNCSIQRGSKKHLALLAPS DVAGWGI FIKDPVQKNEFISEYCGEISQDEAD RRGKVYDKYMC SFLFNLNDFVVD ATRKGNKIRFANHSVNPNCYAKVM MVNGDHRIGIFAKRAIQTGEELFFDY RYSQADALKYVGIEREMEIP	True	False	3.859	5.0	2.622