

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
Q8IXJ9	ASXL1_HUMAN	Homo sapiens	Polycomb group protein ASXL1	35.854505	S199	S501;S503	30397120;21740066;29556021	MKDKQKKKERTWAEARLVLENY SDAPMTPKQILQVIEAEGLKEMRSG TSPLACLNAMLSNSRGGEGLFYKL PGRISLFTLKKDALQWSRHPATVEG EPEPDTADVESCGSNEASTVSGEND VSLDETSNASCSTESQSRPLSNPR DSYRASSQANKQKKKTGVMLPRVVL TPLKVNGAHVESASGFGCHADGES GSPSSSSSGLALGSAAIRQAEVTQ DPAPLLRGRFPATGQMKRNRGEEI DFETPGSILVNTNLRALINSRTFHAL PSHFQQQLLFLPEVDRQVGTDGLL RLSSSALNNEFFTHAAQSWRERLAD GEFTHEMQVRIRQEMEKEKKVEQW KEKFFEDYGGKLGTLKEESLQNV GQEEAEIKSGLCVPGESVRIQRGPAT RQRDGHFKRSRDLRTRARRNLYK KQESEQAGVAKDAKSVASDVPLYKD GEAKTDPAGLSSPHLPGTSSAAPDLE GPEFPVESVASRIQAEPDNLARASAS PDRIPSLPQETVDQEPKQKRKSF QAASASFPEKKPRLEDROQSRNTIES VHTEKPQPTKEEPKVPPIRQLSRIKP PWVVKGOPTYQICPRIIPTTESSCRG WTGARTLADIKARALQVRGARGHHC HREAATTAIGGGGGPGGGGGATDE GGGRGSSSDGGEACGHPEPRGGP STPGKCTSDLQRTQLLPPYPLNGEH TOAGTAMSRARREDLPSLRKEESCL LQRATVGLTDGLGDASQLPVAPTGD QPCQALPLSSQTSVAERLVEQPQL HPDVRTCEESGTTSWESDDEEQGPT VPADNGPIPSLVGDDTLEKGTGQAL DSHPTMKDPVNVTPSSSTPESSPTDC LQNRAFDDELGLGGSCPPMRESDT RQENLKTALVSNSSLHWIPISND EVVKQPKPESREHIPSVEPQVGEW EKAAPTTPALPGDLTAEGLDPLDSL TSLWTVPSRGGSDSNGSYCQVVDIE KLKINGDSEALSPHGESTDTASDFE GHLTEDSSEADTREAAVTKGSSVVK DEKPNWVNSAPLSKVNMDMLVTR TDGMVAPQSWVSRVCAVRQKIPDSL LLASTEYQPRAVCLSMGSSVEATN PLVMQLLQGLSPLKVLPPAHDDSM SESPQVPLTKDQSHGSLRMGSLHGL GKNSGMVDGSSPSSLRALKEPLL SCETGTGLARIEATQAPGAPQKNCK AVPSFDSLHPVTNPITSSRKELEMS KEQFSSFCEDQKEVRAMSQDSNS NAAPGKSPGDLTTSRTPRFSSPNVIS FGPEQTGRALGDQSNVTGQGGKLF GSGNVAATLQRPRPADPMLPAEIPP VFPSGKLGFPSTNSMSGGVQTPRED WAPKPHAFVGSVKNEKTFVGGPLKA NAENRKATGHSPLELVGHLEGMPF VMDLPFWKLPREPGLSEPLEPSS LPSQLSIKQAFYGLSKLQLSSTSFN YSSSPTFPKGLAGSVVQLSHKANF GASHSASLSLQMFDTSSSTVESISLQC ACSLKAMIMCQGGAFCHDDCIGPS KLCVLCVVR	True	False	3.404	5.0	1.949	1.353	1.599	1.638	1.808