

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	extracell region
Q53H47	SETMR_HUMAN	Homo sapiens	Histone-lysine N-methyltransferase SETMAR	24.902576	S441;T442	S508	40596516;38253038	MFAEAAKTRPCGMAEFKEKPEAPT EQLDVACQENLPVGAWPPGAAPAP FQYTPDHVVVPGADIDPTQITFPGCI CVKTPCLPGTCSCLRHGENDNSC LRDIGSGGKYAEPVFECNVLRCRSD HCRNRVVQKGLQFHFQVFKTHKKG WGLRTEFIPKGRFVCEYAGEVLGFS EVQRRIHLOTKSDSNYIIAREHVYN GOVMETFVDPTYIGNIGRFLNHSCE PNLLMIPVRIDSMVPKALFAAKDIV PEEELSYDYSGRYLNLTVSEDKERLD HGKLRKPCYCGAKSCTAFLPFDSSL YCPVEKSNISCGNEKEPSCGGSAPS VFPCKRLTLETMKMMLDKKQIRAI FLFEFKMGRKAAETTRNINNAFGPG TANERTVQWWFKKFKGDESLEDE ERSGRPSEVDNDQLRAIHEADPLTTT REVAEELNVNHSTVVRHLKQIGKVK KLDKWPHELTEHQKRRRFVSSSL ILRNHNEPFLDRIVTCDEKWILYDNR RRSAQWLDQEEAPKHFPKILHPKK VMVTIWWSAAGLIHYSFLNPGETITS EKYAEIDEMNQKLRQLALVNRK GPILLHDNARPHVAQPTLQKLNELG YEVLPHPYSPDLLPTNYHVFKHLN NFLQGKRFFHNQDAENAFQEFVES QSTDFYATGINQLISRWQKCVDCNG SYFD	True	False	2.391	5.0	False	False	1.266	False	