

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
Q53H47	SETMR_HUMAN	Homo sapiens	Histone-lysine N-methyltransferase SETMAR	22.903221	S441;T442	S508	38253038	MFAEAAKTTRPCGMAEFKEKPEAPT EQLDVACGQENLPVGAWPPGAAPAP FQYTPDHVVGPGADIDPTQITFPGCI CVKTPCLPGTCSCLRHGENDNSC LRDIGSGGKYAEPVFECNVLRCRSD HCRNRVVQKGLQFHFQVFKTHKKG WGLRLEFIPKGRFVCEYAGEVLGFS EVQRIHLQTKSDSNYIIAREHVYN GQVMETFVDPTYIGNIGRFLNHSCE PNLLMIPVRIDSMVPKLALFAAKDIV PEEELSYDYSGRYLNLTVSEDKERLD HGKLRKPCYCGAKSCTAFLPFDSSL YCPVEKSNISCGNEKPSMCGSAPS VFPSCKRLTLETMKMMLDKKQIRAI FLFEFKMGRKAAETTRNINNAFPGG TANERTVQWWFKKFKGDESLEDE ERSGRPSEVDNDQLRAIIEADPLTTT REVAEELNVNHSTVVRHLKQIGKVK KLDKWWPHELTENQKNRRFEVSSSL ILRNHNPPFLDRIVTCDEKWILYDNR RRSAQWLDQEEAPKHFPKILHPKK VMVTIWWSAAGLIHYSFLNPGETITS EKYAEIDEMNQKQLRQLALVNRK GPILLHDNARPHVAQPTLQKLNELG YEVLPHPYSPDLLPTNYHVFKHLN NFLQGKRFHNQQDAENAFQEFVES QSTDFYATGINQLISRWQKVCDCNG SYFD	True	False	3.123	5.0	1.621	1.167	0.66	2.328	1.547