

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q96KQ7	EHMT2_HUMAN	Homo sapiens	Histone-lysine N-methyltransferase EHMT2	6.287639	NaN	S40;T44;S47;S140;S173;S232;S242;S246;S350;S412;S413;T555;S569;S1204;T1210	34019948	MAAAAGAAAAAAEGEAPAEMGALL LEKETRGATERVHGSLGDTPRSEET LPKATPDSLEPAGPSSPASVTVTVG EGADTPVGATPLIGDESENLEGDGD LRGGRILLGHATKSFSSPSKGGSCP SRAKMSMTGAGKSPPSVQSLAMRL LSMPGAQGAAAAGSEPPATTSP QPKVHRARKTMSKPGNGQPPVPEK RPPEIQHFRMSDDVHSLGKVTSDLA KRRKLNSSGGGLSEELGSARRSGEVT LTKGDPGSLEEWETVVGDDFSLYYD SYSVDERVSDSKSEVEALTEQLSE EEEEEEEEEEEEEEEEEEEEEEEEDE ESGNQSDRSGSSGRRKAKKKWRKD SPWVKPSRKRKRREPPRAKEPRGVN GVGSSGPSEYMEVPLGSLELPSEGT LSPNHAGVSNDTSSLETERGFEELP LCSCRMEAPKIDRISERAGHKCMAT ESVDGELSGCNAAILKRETMRPSSR VALMVLCEHRARMVKHHCCPGCG YFCTAGTFLECHPDFRVAHRFHKAC VSQLNGMVFCPHCGEDASEAQEVTI PRGDGVTTPAGTAAPAPPPLSQDVP GRADTSQPSARMRGHGEP RRPPCD PLADTIDSSGPSLTLPNGGCLSAVGL PLGPGREALEKALVIQESERRKLR F HPRQLYLSVKQGELOKQVILMLLDNL DPNFQSDQQSKRTPHAAAQKGSVE ICHVLLQAGANINAVDKQQRTP LME AVVNNHLEVARYMVQRGGCVSKE EDGSTCLHHAAKIGNLEMVSL LLST GQVDVNAQDSGGWTPIIWAAEHKHI EVIRMLLTRGADVTLTDNEENIC LH WASFTGSAAIAEVLLNARCDLH AVN YHGDTPHIAARESYHDCVLLFLSR GANPELRNKEGDTAWDLTPERSDV WFALQLNRKLRG VGNRAIRTEKIIC RDVARGYENVPIPCVNGVDGEP CPE DYKYISENCETSTMNIDRNITHLQ H CTCVDDCSSSNCLCGQLSIR CWYDK DGRLLQEFNKIEPPLIFECNQAC SC WRNCKNRVVQSGIKVRLQLYRTAK MGWGVRALQTIPOGTFICEYV GELIS DAEADVREDDSYLFDLDNKDGE VYC IDARYYGNISRFINHLCDPN IIPVRVF MLHQDLRFPRIAFFSSRD IRTGEELG FDYGDRFWDIKSKYFTCQCG SEKCK HSAEAIQSRRLARLD PHPELLPEL GSLPPVNT

