

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q12830	BPTF_HUMAN	Homo sapiens	Nucleosome-remodeling factor subunit BPTF	53.72858	S1617;S1626;T1709;T1712;S1742;S1749;T1750;T1753;T1754;T1757;T1759;S2081;S2083;T2094;T2103;S2140;T2148;T2192;S2229;T2230;S2233;T2237;S2239;S2240;T2241;T2346;S2348	S216;S572;S763;S817;T1064;S1231;S1300;T1303;S1310;S2098;S2465	30379171;33214551;31492838;32119511;30620550;31637018;20068230;30059200;29249144;28657654;29351928;29237092;21158410;34019948;3626323;27114449;20305658;22661428;32574038;34065225	MRGRRGRPPKQPAAPAAERCAPAPP PPPPPTSGPIGGLRSRHRGSSRRGR WAAQAQAEVAPKTRLSFRGGSSRRR KPPPPPPAPPSTAPGRGGRRGGGGG RTGGGGGGGHLARTTAARRAVNKV VYDDHESEEEEEEDMVSEEEEEEE DGDAEETQDSEDEEDEEDEEDDDDD SDYPEEMEDDDDDASYCTESSFRSH STYSSTPGRRKPRVHRPRSPILEEKD IPPLEFPKSEEDLMVPNEHIMNVIAI YEVLRNFGTVLRLSPFRFEDFCAALV SQEQCTLMAEMHVLLKAVLREED TSNTTFGPADLKDSVNSTLYFIDGM TWPEVLRVYCESDKEYHHVLPYQEA EDYPYGPVENKIKVLQFLVDQFLTTN IAREELMSEGVIQYDDHCRVCHKLG DLLCCECTCSAVYHLECVKPPLEEVPE DEWQCEVCVAHKVPGVTDCAEIQK NKPYIRHEPIGYDRSRRKYWFLNRR LIIEDTENENEKKIWYYSTKVQLAE LIDCLDKDYWEAELCKILEEMREEIH RHMDITEDLTNKARGSNKSFLAAAN EEILESIRAKKGDIDNVKSPEETEKD KNETENDSKDAEKNREEFEDQSLSE KDSDDKTPDDDDPEQKGSEEPTEVGD KGNVSANLGDNTTATSEETSPSE GRSPVGCSETPDSSNMAEKKVASE LPQDVPEEPNKTCESSNTSATTTSIQ PNLENSNSSSELNSSQSESAKAADD PENGERSHTPVSIQEEIVGDFKSEK SNGELSESPGAGKGASGSTRITRLR NPDSKLSQLKSQQVAAAHEANKLF KEGKVLVNSQGEISRLSTKKEVIM KGNINNYFKLQGEQKYRVYHNQYST NSFALNKHQHREDHDKRRHLAHKF CLTPAGEFKWNGSVHGSKVLITISTL RLTITQLENNIPSSFLHPNASHRA NWKAVQMCKPREFALALALECA VKPVVMLPIWRESLGHTRLHRMTSI EREKEKVKKKEKKQEEEEETMQQA TWVKYTFPVKHQVWKQKGEEYRVT GYGGWSWISKTHVYRFVPKLPNGTN VNYRKSLEGTKNMMDENMDESDK RKCSRSPKKIKIEPDSEKDEVKGSDA AKGADQNEMDISKITEKKDQDVKEL LDSDSKPKKEEPMEVDDDMKTES HVNCQESSQVDVNVVSEGFHLRTSY KKKTKSSKLDGLLERRIKQFTLEEKQ RLEKIKLEGGIKGIGKTSTNSSKNLS ESPVITKAKEGCQSDSMRQEQSPNA NNDQPEDLIQGCSESDSSVLRMSDP SHTTNKLYPKDRVLDVVSIRSPETKC PKQNSIENDIEKVSDLASRGQEPSK SKTKGNDFIDDSKLASADDIGTLIC KNKKPLIQEESDTIVSSSKSALHSSV PKSTNDRDATPLSRAMD FEGKLGCD SESNSTLENSSDTVSIQDSSEEDMIV QNSNESISEQFRTRREQDVEVLEPLK CELVSGESTGNCEDRLPVKGTEANG KKPSQQKLEERPVNKCSAQIKLKN TTDKNNENRESEKKGQRTSTFQIN GKDNKPKIYLKGECLKEISESRVVS NVEPKVNNINKIIPENDIKSLTVKES

AIRPFINGDVIMEDFNERNSSSETKS
HLLSSDAEGNYRDSLETLPSTKES
DSTQTTTPSASCPESENSVNQVEDME
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KLQSTAS