

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P00519	ABL1_HUMAN	Homo sapiens	Tyrosine-protein kinase ABL1	15.414137	S840;S855	S50;T70;T115;T128;T139;T172;T185;T215;T226;S229;T253;T257;T393;T413;S446;S559;S569;S618;S619;S620;S659;S683;S718;T735;T751;T781;T814;T823;T844;T852;S855;S917;S977	30059200;35254053;28657654	MLEICLKLVGCKSKKGLSSSSSSCYLE EALQRPVASFDFEPQGLSEAARWNSK ENLLAGPSENDPNLFVALYDFVASG DNTLSITKGEKLRVLGYNHNGEWCE AQTKNGQGWWPSNYITPVNSLEKHS WYHGPPVSRNAAEYLLSSGINGSFLV RESESSPGQRSISLRYEGRVYHYRIN TASDGKLYVSSSRFNTLAELVHHH STVADGLITTLHYAPKRNKPTVYGV SPNYDKWEMERTDITMKHKLGGGQ YGEVYEGVWKKYSLTVAVKTLKEDT MEVEEFLKEAAVMKEIKHPNLVQLL GVCTREPPFYIITEFMTYGNLLDYLR ECNRQEVNAVLLYMATQISSAMEY LEKKNFIHRDLAARNCLVGENHLVK VADFGLSRLMTGDTYTAHAGAKFPI KWTAPESLAYNKFSIKSDVWAFGVL LWEIATYGMSPYPGIDLSQVYELLEK DYRMERPEGCPEKVYELMRACWQW NPSDRPSFAEIHQAFETMFQESSISD EVEKELGKQGVRGAVSTLLQAPELP TKTRTSRRAAEHRDTTDVPMPHSK GQGESDPLDHEPAVSPLLPRKERGP PEGGLNEDERLLPKDKKTNLFSALIK KKKKTAPTTPPKRSSSFREMDGQPER RGAGEEEGRDISNGALFTPLDTAD PAKSPKPSNGAGVPNGALRESGGSG FRSPHLWKKSSTLTSSRLATGEEEG GGSSSKRFLRSCSASCVPHGAKDTE WRSVTLPRDLQSTGRQFDSSTFGGH KSEKPALPRKRAGENRSDQVTRGTV TPPPRLVKKNEEADEVFKDIMESS PGSSPPNLTpkplrrqvtvapasglp HKEEAGKGSALGTPAAAEPVTPTSK AGSGAPGGTSGKPAEESRVRHKHS SESPGRDKGKLSRLKPAPPPPAASA GKAGGKPSQSPSQEAAGEAVLGAKT KATSLVDVAVNSDAAKPSQPGGLK PVLPATPKPQSAKPSGTPISPAPVPST LPSASSALAGDQPSSTAFIPLISTRVS LRKTRQPPERIASGAIKGVVLDSTE ALCLAISRNSEQMASHSAVLEAGKN LYTFCVSYVDSIQQMRNKFAFREAIN KLENNLRELQICPATAGSGPAATQD FSKLLSSVKEISDIVQR